Command line parameters:

MODEL-frame+\_p2n.model -DEV-xlp
-Q-/cgn2\_1/USPT0\_spool/US09801115/runat\_20062003\_141103\_13596/app\_query.fasta\_1.263
-DB-EST -QFMT-fastap -SUFFIX-rst -MINNATCH-0.1 -LOOPCL-0 -LOOPEXT-0
-UNITS-bits -START-1 -END-1 -MATRIX-blosum62 -TRANS-human40.cdi -LIST-45
-DCCALIGN-200 -THR\_SCORE-pct -THR\_MAX-100 -THR\_MIN-0 -ALIGN-15 -MODE-LOCAL
-OUTFMT-pto -NORM-ext -HEAPSIZE-500 -MILEN-0 -MAXLEN-2000000000
-USER-US09801115\_eCGN\_1\_1\_2874\_@runat\_20062003\_141103\_13596 -NCPU-6 -ICPU-3
-NO\_MMAP -LARGEQUERY -NEG\_SCORES-0 -WAIT -DSPBLOCK-100 -LONGLOG
-DEV\_TIMEOUT-120 -WARN\_TIMEOUT-30 -THREADS-1 -XGAPOP-10 -XGAPEXT-0.5 -FGAPEXT-7
-FGAPEXT-7 -YGAPOP-10 -YGAPEXT-0.5 -DELEXT-7 Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries OM protein - nucleic search, using frame\_plus\_p2n model Database : Minimum DB seq length: 0
Maximum DB seq length: 2000000000 Total number of hits satisfying chosen parameters: Scoring table: Sequence: Perfect score: Run on: June 25, 2003, 12:44:54; Search time 1420 Seconds (without alignments) 1129.122 Million cell updates/sec Xgapop 10.0 , Xgapext Ygapop 10.0 , Ygapext Fgapop 6.0 , Fgapext Delop 6.0 , Delext BLOSUM62 508 10: 11: 12: 13: 14: 15: 16: 16: 17: 17: 18: 19: 20: 21: 22: 23: 24: 25: 16154066 seqs, 8097743376 residues US-09-801-115B-2 MDNVQPKIKHRPFCFSVKGH.......FNPSGPYQKKPVHEKKEVL GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd. em\_estmu:\*
em\_estov:\* em\_estom:\*
gb\_gss:\*
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em\_gss\_inv:\*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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### ALIGNMENTS

REFERENCE AUTHORS TITLE		SOURCE ORGANISM	KEYWORDS	VERSION	ACCESSION		NOTE	LOCUS	AA911088/c	RESULT 1
<pre>1 (bases 1 to 386) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),</pre>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	human. Homo sapiens	EST.	AA911088.1 GI:3050378	AA911088	MONA COCIONACIO	al NCT CGAP GC4 Homo sapiens cDNA	AA911088 386 bp mRNA linear EST 09-JUN-1998		·

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Insert Length: 381 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
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human
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
Emmert-Buck, M.D., Ph.D.
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/clone_lib="NCI_CGAP_GC4"
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/lab_host="DH10B"
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AW183193 396 bp mRNA linear ES: xj67f11.xl Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone
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Contact: Robert Strausberg, Ph.D.
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disease"
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                                                                                                                                                                     61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                                                                                                            41 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe 60
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP).
AA516431
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                                                                                                                                     GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGCCCTTATTTACCGGAAGCTTCTG
                                                                                                                                                                                                         GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                                                                                                                                                                                                                                                  ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-30287, 682632-687239, 726408-728711, and 72996-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
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/clone_lib="Soares_NFL_T_GBC_S1"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                        21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
81 PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
                                                                                                61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
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Insert Length: 527 Std Error: 0.00
Seg primer: -40ml3 fwd. ET from Amersha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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/db_xref="taxon:9606"
/clone="IMAGE:901516"
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/lab_host="DH10B"
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Email: cgapbs-r@mail.nih.gov

This clone is available royalty-free through LINL; contact the

IMAGE Consortium (info@image.llnl.gov) for further information.

Seg primer: -40Up from Gibco.
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Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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402 bp mRNA linear EST 27-OCT-19
WU21109.xl Soares_Dieckgraefe_colon_NHCD Homo sapiens cDNA clone
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                                         ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyValPhe 60
                                                                                 GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
       GTATCTGTGTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT 189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                went through one round of normalization. Tissue samples provided by Dr. Brian Dieckgraefe (Washington University, dieck@im.wustl.edu); colonic mucosa represents a range of disease involvement from moderate to severe Crohn's disease; samples include both perforating (fistulas) and non-perforating samples. Library constructed by Bento Soares and M. Fatima Bonaldo. "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the
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/db_xref="taxon:9606"
/clone="taxon:9713"
/clone=lib="Soares_Dieckgraefe_colon_NHCD"
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BM553628
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     152
21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
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                                                                                                                                                                        1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
http://image.llnl.gov
http://image.llnl.gov
http://image.llnl.gov
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http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 404)
NIH-MCC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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BM553628
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                           GTGAAGATGCTGCGGCTGGATATTATTAACTCACTGGTAACAACAGTATTCATGCTCATC
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                                                                                                                                                                                                                                                                                                                                                                                                                                    /Clone_lib="NIH_MGC_88"
/tissue_type="duodenal adenocarcinoma, cell line"
/lab_host="DH108 (phage-resistant)"
/note="Organ: small intestine; Vector: pCMV-SPORT8;
Site_1: Not1; Site_2: Sall; Cloned unidirectionally;
oligo-dT primed. Average insert size 1.767 kb. Library
enriched for full-length clones and so structed by Life
Technologies. Note: this is a NIH_MGC Library."
a 80 c 87 g 116 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5740478"
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                                                                                                                                US-09-801-115B-2 (1-99) x AI826623 (1-409)
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Best Local Similarity:
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                   21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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wk35d10.x1 NCI_CGAP_Pr22 Homo
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
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Seq primer: -40UP from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
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                                                                                                                                                                                                                                                                                                                                    /note="Organ: prostate; Vector: pT/T3D-Pac (Pharmacia)
with a modified polylinker; 1st strand cDNA was prepared
from normal prostate bulk tissue, and was then primed with
a Not I - oligo(dT) primer. Double-stranded cDNA was
ligated to Eco RI adaptors (Pharmacia), digested with Not
I and cloned into the Not I and Eco RI sites of the
modified pT/T3 vector. Library is normalized, and was
constructed by Bento Soares and M. Fatima Bonaldo. "

80 g 123 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2417395"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="NCI_CGAP_Pr22"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="normal prostate"
/lab_host="DH10B"
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US-09-801-115B-2 (1-99) x AI128804 (1-423)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: -40m13 fwd. ET from Amersham High quality sequence stop: 413.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                              NbHL19W."
                                                                                                                                                                                                                                                                                                                                                    normalization to a Cot = 5. Library constructed by M.Fatima Bonaldo. This library was constructed from the
                                                                                                                                                                                                                                                                               double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified p773 vector (Pharmacia). Library went through one round of
                                                                                                                                                                                                                               same fetus as the fetal lung library, Soares fetal lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="IMAGE:1694391"
/clone_1ib="Soares_fetal_heart_NbHH19W"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /sex="unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                             dev_stage="19 weeks"
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1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -41ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R. Wash Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 314 286 1800
Fax: 314 286 1810
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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                                                              /note-"Organ: mixed (see below); Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_1: Ecor RI; Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479."
                                                                                                                                                                                                                                                                                                                               /tissue_type="pooled human melanocyte, fetal heart, pregnant uterus"
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/clone="IMAGE:812244"
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                       Email: estewatson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -41ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                          Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1810
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 432)
Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S.,
Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B.
Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylle
,T., Waterston,R. and Wilson,R.
Washbr-Merck EST Project 1997
Unpublished (1997)
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AA429945
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zw67f10.sl Soares_testis_NHT Homo sapiens cDNA clone
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Location/Qualifiers
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/organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                   /sex="male"
                                                                                                                                                                                    /clone_lib="Soares_testis_NHT"
                                                                                                                                                                                                           /clone="IMAGE:781291"
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0237h05 xl Soares_NhHMPu_Sl Homo sapiens 3', mRNA sequence.
                                                                                                                                                                                                                                                                                         Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 452)
                                                                                                                                                                                                                                                                                                                                               Tumor Gene Index
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                             IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                             This clone is available royalty-free through LLNL ; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
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          /note-"Organ: mixed (see below); Vector: pT/T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus)
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                                                                                              pregnant uterus"
                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1677561"
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                                                                                                             /tissue_type="Pooled human melanocyte, fetal heart,
                                                                                                                               /clone_lib="Soares_NhHMPu_S1"
                                                                                                                                                                                                            Location/Qualifiers
                                                                             /lab_host="DH10B"
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453 bp mRNA linear EST 20-DEC wg90a02.xl Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE:2372330 3', mRNA sequence.
                                                                                                                                                                                                     Unpublished (1997)
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National Cancer Institute, Cancer Genome Anat
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AI743235.1 GI:5111523
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/lab_host="DH10B"
/lab_host="DH10B"
/note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from five normalized
                                                                             /clone="IMAGE:2372330"
/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
                                                                                                                          /db_xref="taxon:9606"
                                                                                                                                            /organism="Homo sapiens"
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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468 bp mRNA linear EST 20-OCT-2000 7171f03.xl Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone IMAGE:3526805 3' similar to TR:Q9UI41 Q9UI41 CHEMOKINE-LIKE FACTOR
                                                                                                                                                                                                                                                                                                                                                                                     BF109912
                                                                                        This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Seq primer: -40UP from Gibco.
                                                                                                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                    Tumor Gene Index
Unpublished (1997)
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309384-310919, 323208-325895 Soares NbHSF pool 1:

145032-147335, 147720-148103, 148872-149255, 15002 -
150407, 151176-152327 Soares Nb2HF8-9W pool 1:

758280-760583, 772104-774407 Soares NbHPA pool 1:
304776-306311, 320136-322823, 326280-32663 Soares NbHOT pool 1: 723720-724407, 739080-740999 Subtraction by Bento Soares and M. Fatima Bonaldo."
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3526805"
                                                                       Location/Qualifiers
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US-09-801-115B-2 (1-99) x BF109912 (1-468)
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Rattus norvegicus
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UI-R-CA1-bjb-b-12-0-UI.sl UI-R-CA1 Rattus norvegicus cDNA clone
UI-R-CA1-bjb-b-12-0-UI 3', mRNA sequence.
                    Contact: Soares, MB
Program for Rat Gene
                                                                  97044477
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                                                                                    Genome Res. 6 (9), 791-806 (1996)
                                                                                                             discovery
                                                                                                                                                  Bonaldo, M.F.,
                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
University of Iowa
                                                                                                                                 Normalization and subtraction:
                                                                                                                                                                                               Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AlaLeuValThrAlaValCysCysLeuAlaAspClyAlaLeuIleTyrArgLysLeuLeu 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MetAspAsnValGlnProLysIleLysH1sArgProPheCysPheSerValLysGlyH1s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
                                                                                                                                                                         (bases 1 to 510)
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                                                                                                                                                  Lennon, G. and Soares, M.B.
                        Discovery and
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RESULT 15
BG705303
                                                                                                                      DEFINITION
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                                     KEYWORDS
                                                            VERSION
                                                                              ACCESSION
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                                     BG705303
BG705303.1 GI:13979504
EST.
                                                                                                     BG705303 524 bp mRNA linear EST U/-MAY-2 602687808F1 NIH_MGC_95 Homo sapiens cDNA clone IMAGE:4820568 5',
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: msoares@blue.weeg.uiowa.edu
The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. cDNA Library Preparation: M.B. Soares Lab Clone distribution: clones will be available through Research Genetics (www.resgen.com)
                                                                                             mRNA sequence
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Tel: 319 335 8250
Fax: 319 335 9565
                    numan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="DH10B (life Technologies)"
/note="Vector: pT773D-Pac (Pharmacia) with a modified
polylinker; Site_1: Not I; Site_2: Eco RI; The UI-R-CA1
library is a subtracted library derived from the following
tissues: thalamus, cerebellum, hypothalamus, medulla, pons
, midbrain, cerebral cortex, corpus striatum, testis, and
hippocampus. For a detailed description of the library
from which this clone was derived, please visit our web
site at ratest-eng-ulowa.edu. The subtraction has been
previously described in (Bonaldo, Lennon and Soares,
Genome Research 6:791-806, 1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TAG_SEQ=None found"
131 c 104 g
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/strain="Sprague-Dawley"
/db_xref="taxon:10116"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="UI-R-CA1"
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          381
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61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                                                                                                                                                                201 GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                                                                                                                            41 ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 524)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 521.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     http://image.llnl.gov
Plate: LLAM10725 row: o column: 01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4820568"
/clone_lib="NIH_MGC_95"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="hippocampus"
/lab_host="DH10B"
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Maximum DB seq length: 2000000000
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-Q-/cgn2_1/USPTO_spoo1/USO9801115/runat_20062003_141102_13575/app_query.fasta_1.263

-DB-M_Genneseq_101002 -QFMY-fastap -SUFFIX-ring -MINMATCH=0.1 -LOOPCL=0

-LOOPEXT=0 -UNITS-bits -START-1 -END--1 -MATRIX-blosum62 -TRANS-human40.cdi

-LIST-45 -DOCALIGN-200 -THR_SCORE-pct -HAAX-100 -THR_MIN=0 -ALIGN-15

-MODE-LOCAL -OUTFMY-pto -NORM-ext -HEAPSIZE_500 -MIXLEN=0 -MAXLEN-200000000

-USER-USO9801115_@CGN_1_1_396_@runat_20062003_141102_13575 -NCPU=6 -ICPU=3

-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -MAIT -DSPBLOCK=100 -LONGLOG

-DEV_TIMEOUT=120 -MARN_TIMEOUT=30 -THEADS-1 -XGAPDEXT=0.5 -FGAPOP=6

-DEV_TIMEOUT=120 -MARN_TIMEOUT=30 -THEADS-1 -XGAPDEXT=0.5 -FGAPOP=6
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1987.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1988.DAT:*
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                                /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1998.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1999.DAT:*
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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2001B.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

### SUMMARIES

	c c c 39		33222222 344 340	116 17 18 19 20 21 23	6 C C 9 110 112 133	Result No. 1 2 3 3
215 215 211 211	245 245	318.5 318.5 318.5 317 294 257.5	467.5 461 461 357 339.5 339.5 339.5 331 323	471.5 471.5 471.5 471.5 471.5 471.5 471.5	508 508 508 508 508 508 504 471,5	Score 508 508 508 508
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AAK68359 AAK68360 AAK68361 AAK82406	ABL62782 ABL62959 ABL63180	AAZ 34052 AAC 78499 AAF 93358 AAX 41509 AAA 41512 ABA 09082 ABA 09082	ABK35884 AAC00147 AAA16090 AAA38009 AAS45120 AAS45121 AAA38008 AAX97651	AAZ56747 AAZ34051 AAC78498 AAC58238 AAA67771 AAF64053 ABL90366 AAS44933	AAF64012 AAA38006 AAH98515 AAH98548 AAH344835 AAS44932 AAAA15919 AAAA15919 AAA377826 AAA397826	ID 
immune/ immune/ immune/ immune/ immune/	t cancer	Human EST DNA43509 Human EST DNA43509 cDNA encoding SRT Human secreted pro Human secreted exp Human transmembran Gene #744 used to	sequent secuent secuen	n transmembra n PRO772 nucl n PRO772 (UNC) n PRO772 nucl n PRO772 nucl n secreted pr encoding hum n polynucleot encoding nov	encoding handleotic EST-deriv EST-deriv colon car encoding r protein c secreted nucleotic	Description  Human secreted pro cDNA sequence #397  Human secreted pro Human secreted pro

### ALIGNMENTS

# RESULT 1 AAX97873

AAX97873 standard; cDNA; 413 BP.

# AAX97873;

23-SEP-1999 (first entry)

Human secreted protein encoding cDNA #61.

Secreted protein; human; cytostatic; thrombotic; osteopathic; forensic; diagnostic; gene therapy; chromosome mapping; secretion vector; ss.

# Homo sapiens

WO9925825-A2

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13-NOV-1997;
17-DEC-1997;
09-FEB-1998;
13-APR-1998;
10-AUG-1998;
immune
 Human secreted protein; hyperproliferative disorder; autoimmune disorder;
immune deficiency disorder; blood disorder; inflammatory disorder;
                                 cDNA sequence #397 encoding novel human secreted protein.
                                                          08-MAY-2002
                                                                                   ABK36006
                                                                                                          ABK36006 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX97813-X97906 represent extended cDNA's which encode novel human secreted proteins (see AAY36129-Y36222) and which have cytostatic, thrombotic and osteopathic activity. The extended cDNAs can be used to express secreted proteins or parts of them or to obtain antibodies capable of binding to the secreted proteins. They may also be used in diagnostic, forensic, gene therapy and chromosome mapping procedures. Uses also include design of expression vectors and secretion vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 413 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Extended cDNAs encoding secreted proteins
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                                                                                                                                                                                                                                                                                                                                               ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                                                                                                                                                                                                                                ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
                                                                                                                                                                    PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu
                                                                                                                                                                                                                   AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
                                                                                                                                                                                                                                                                                                   GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                                                                                                       GCACTTGTGACAGCAGTATGCTGCCTTGCCGACGGGCCCCTTATTTACCGGAAGCTTCTG
                                                                                                                                                                                                                                                     GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 261;
                                                          (first entry)
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98US-0074121.
98US-0081563.
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61

AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu

80

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200

260

GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT ValSerValLeuAlaLeuIleProGluThrThrThrThrLeuThrValGlyGlyGlyValPhe 141

21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle

40

20

ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC

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US-09-801-115B-2 (1-99) x ABK36006
                                                                                                Query Match:
                                                                                                                Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                  Alignment Scores:
                                                                                                                                                                                                                                           The present invention relates to the isolation of novel cDNA sequences which encode human secreted proteins. The cDNA sequences have been derived from a variety of human tissues. The invention also provides a method for producing proteins from these polynucleotide sequences. The proteins are useful for identifying compounds that modulate their activity and production. The sequences of the invention are useful for treating diseases such as hyperproliferative disorders (e.g. cancer), immune deficiency disorders (e.g. severe combined immunodeficiency (SCID)), autoimmune disorders (e.g. multiple sclerosis), blood disorders (e.g. thrombocytopaenia), inflammatory disorders (e.g. arthritis), infectious disorders (e.g. hepatitis), allergic conditions (e.g. asthma), neurodegenerative disorders (e.g. Alzheimer's disease), liver fibrosis, coagulation disorders (e.g. haemophilia), and tumours. The polynucleotide sequences of the invention are also useful in gene therapy. ABK35610-ABK36232 represent the cDNA sequences of the invention that encode for novel human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Merberg D
Clark HF,
                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Six hundred and twenty three polynucleotides derived from a variety of human tissue sources which encode secreted proteins, useful for treating immune deficiencies and disorders such as autoimmune disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 296-297; 393pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-179322/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infectious disorder; allergic condition; neurodegenerative disorder;
liver fibrosis; coagulation disorder; gene therapy; antimicrobial;
tumour; cancer; hepatotropic; immunosuppressive; antirheumatic; gene
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1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
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                                                                                                                                                                                                                  96
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                                                (1-439)
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lding V, Wong
ca K, Graham ;
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RESULT 3
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07-MAR-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzhelmer's disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthittis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
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970S-0043311.
970S-0043313.
970S-0043356.
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23-MAY-1997;
                                                                                                                                                             Feng
Kyaw
  disorders, immune diseases, inflammation
    New isolated human genes and the secreted polypeptide(s) they encode - useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
                                                                               WPI; 1998-506364/43.
P-PSDB; AAW74818.
                                                                                                                                          Ruben
                                                                                                                                                                                               Bednarik DP,
                                                                                                                                                                                                                                        (HUMA-) HUMAN
                                                                                                                                    arik DP, Brewer LA, Carter KC, Duan R, Ebner R, I
P, Ferrie AM, Fischer CL, Florence KA, Greene JM
H, Lafleur DW, Li Y, Moore PA, Ni J, Olsen HS,
n SM, Shi Y, Soppet DR, Young PE, Yu GL, Zeng Z;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC This sequence represents a nucleic acid molecule designated Gene 88 from CC the human cDNA clone HADAV32 (deposited as clone ATCC 97901 and ATCC C2 09047) which encodes a secreted human protein. The gene can be used to generate fusion proteins by linking to the gene to a human CL immunoglobulin Fc portion (e.g. AAV59502) for increasing the stability of the fused protein as compared to the human protein only.

CC The invention relates to 186 novel genes and their fragments (nucleic cacid sequences: AAV59511-V59812; amino acid sequences AAW74731-W75026) CC which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypeptides in a CL sample or by determining the presence of mutations in the new CC can be diagnosed by determining the amount of the new polypeptides in a CL sample or by determining the described for each of the 186 CC polynucleotides. Specific uses are described for each of the 186 CC polynucleotides, based on which tissues they are most highly expressed in (see AAV59511 for described uses).
                                                 Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restencis; AIDS; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
        WO9839448-A2
                                                                                                                                                                                                19-JAN-1999 (first entry)
                                Homo sapiens
                                                                                                                                                                         Human secreted protein gene 88 clone HAUAV32.
                                                                                                                                                                                                                                                      AAV59746 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 504 BP;
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                                                                                                                                                                                                                                                                                                   AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                                                                                                                                                                                                                                                                                    ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
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    08-JUL-1997
16-JUL-1997
18-AUG-1997
22-AUG-1997
22-AUG-1997
22-AUG-1997
22-AUG-1997
22-AUG-1997
22-AUG-1997
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11-APR-1997
23-MAY-1997
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07-MAR-1997;
07-MAR-1997;
07-MAR-1997;
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13-JUN-1997;
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11-APR-1997
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07-MAR-1997;
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97US-0040336.
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Percent Similarity:
Best Local Similarity:
Query Match:
DB:
                                                 Score:
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                                                                                                                                  This sequence represents a nucleic acid molecule designated Gene 88 from the human cDNA clone HAUAV32 (deposited as clone ATCC 97897 and ATCC 209043) which encodes a secreted human protein. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin Fc portion (e.g. AAV55902) for increasing the stability of the fused protein as compared to the human protein only.

The invention relates to 186 novel genes and their fragments (nucleic acid sequences: AAV59511-V59812; amino acid sequences AAV74731-W7526) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypucleotides. Specific uses are described for each of the 186 polypucleotides, based on which tissues they are most highly expressed in
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Feng
Kyaw
                                                                                                                                                                                                                                                                                                                                 Claim 1;
                                                                                                                                                                                                                                                                                                                                                       New isolated human genes and the secreted polypeptide(s) they encode - useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bednarik
                                                                                                   Sequence 506 BP;
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Ferrie AM, Eischer CL, Florenus
Flair DW, Li Y, Moore PA, Ni
                                                                                                                                                                                                                                                                                                                              Page 472; 721pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENOME SCI INC
                                                                                                                           for described uses).
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97US-0056881.
97US-0056882.
97US-0056884.
97US-0056886.
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97US-0057761.
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97US-0056903.
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970S-0056879.
                                                                                                  148 A; 103 C;
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Matches:
                       Mismatches:
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    This sequence represents human cDNA encoding a secreted protein. The invention relates to sequences AAA87725-A8774 which encode human secreted proteins AAB25763-B25812. The proteins include signal peptides. Included in the invention are a host cell containing one of the cDNA sequences, and a purified antibody capable of binding to one of the secreted proteins. Also contained in the invention are methods for storing the sequence data on a computer system, and a method for identifying features of the cDNA sequences using a computer programme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               mitochondriocytopathy; diabetes; atherosclerosis; Alzheimer's neurodegenerative disorder; graft rejection; dementia; hyperl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; secreted protein; forensic procedure; gene therapy; chromosome mapping; cancer; autoimmune disease; cardiovascular disord cystic fibrosis; hypothyroidism; immunological disorder; amyloidosis; brain disorder; skeletal muscle disorder; eye disorder; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-NOV-2000
                                                                                                                       Claim 1; Page 169-170; 306pp; English.
                                                                                                                                                                Polynucleotides and peptides, useful in
                                                                                                                                                                                                                                                                                                   22-DEC-1998;
25-JUN-1999;
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                                                                                                                                                  mapping procedures
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    shock;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    impotence;
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                                                                                                                                                                                                                                               Dumas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA;
                                                                                                                                                               polypeptides encoding proteins with signal diagnostic, forensic, gene therapy and chr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           encoding cDNA SEQ ID #29.
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 WPI;
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                       Dumas Milne Edwards
                                                                     25-JUN-1999;
21-DEC-1999;
                                                                                                                                04-JAN-2001
                                                                                                                                                                                                                                     cDNA encoding human
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                                                                                                                                                                                                                                                                                      AAF64012
                                               (GEST ) GENSET
                                                                                                         21-JUN-2000;
                                                                                                                                                      WO200100806-A2
                                                                                                                                                                               Homo
                                                                                                                                                                                                              Secreted protein; prevention; treatment; diagnosis; disease;
                                                                                                                                                                                                                                                                                                           AAF64012 standard;
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                                                                                                                                                                                                     infection;
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 2001-071487/08
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RESULT 7
AAA38006
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DB:
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Best Local Similarity:
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                                                WPI; 2000-388170/34
                          P-PSDB; AAY98142
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                                                                                           Ma D,
                                                                                                                                                                                                                                   14-MAY-1999;
                                                                                                                                                                                                                                                                                                                                 CN1244584-A
                                                                                                                                                                                                                                                                                                                                                                                                                      radiotherapy;
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                                                                                                                                         (UYBE-) UNIV
                                                                                                                                                                                       14-MAY-1999;
                                                                                                                                                                                                                                                                                 16-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                 UCK-1; chemotaxis factor; immunocyte; haemopoiesis stimulant; tumour;
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                                                                                           Han W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu
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                                                                                                                                         BEIJING MEDICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                      chemotherapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein expression.
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Matches:
Conservative:
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RESULT 8
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Best Local Similarity:
Query Match:
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25-JAN-2000;
17-JUL-2000;
03-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents an UCK-1 cDNA sequence encoding a chemotaxis factor polypeptide. The UCK-1 protein exhibits immunocyte chemotaxis activity and a haemopoiesis stimulating effect. The invention relates t UCK proteins, their encoding nucleotide sequences and antibodies and antagonists against the proteins. The nucleotide and protein sequences are useful for the preparation of a composition for the diagnosis and treatment of diseases associated with abnormal immunocyte function and low haemoppies function caused by radiotherapy and chemotherapy used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chemotaxic factor useful for treatment and diagnosis of immunocyte disorders - has immunocyte chemotaxic stimulating factor
                                                                                                                                 tomato; monkey; dog; sea urchin; expressed sequence tag; ES diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition; ss.
                                                                                                                                                                                       Human EST-derived coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 534 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treat tumours and other diseases.
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                                                                                                              Homo
                                                                                                                                                                  Human; sheep;
                                                                                                                                                                                                              12-OCT-2001
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                                            25-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                               AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu 80
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; 2000US-0491404.
; 2000US-0617746.
; 2000US-0631451.
                                                                                                                                                                                                             (first entry)
                                             2001WO-US02687
                                                                                                                                                                 pig;
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                                                                                                                                                                  COW;
                                                                                                                                                                fruit fly; yeast; hamster; macaque;
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Mismatches:
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Matches:
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RESULT 9
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AAC AAH985
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides the protein and coding sequences proteins from a variety of organisms, including human, dog, cat, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, urchin and tomato. These were derived from expressed sequence to from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated polypeptide for antibodies and research
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                                          Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; tomato; monkey; dog; sea urchin; expressed sequence tag; ES' diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition; ss.
                                                                                                                                                                               AAH98548;
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                                                                                                                   EST-derived coding sequence
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                                                                                                                                                                                                                                                                                                                                              AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                                                                                                                                                                                                                                                                           ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATC
                                                                                                                                                                                                            standard;
                                                                                                                                                                                                                                                                                      PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu
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Homo sapiens

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17-JUL-2000;
03-AUG-2000;
           03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, pig, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags (ESTs) from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a cDNA of the contraction.
                                AAH34835
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the invention.
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                                                                                                          81 PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
                                                                                                                                                        61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
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                                                                                                                                                                                       GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
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                                                                                               Liu C,
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2000US-0617746.
2000US-0631451.
2000US-0663870.
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US-09-801-115B-2 (1-99) x AAH34835 (1-558)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon CC cancer associated nucleic acid molecules (N) and proteins (P), where CC the proteins are collectively known as colon cancer antigens. The colon CC cancer antigens have cytostatic activity and can be used in gene CC therapy and vaccine production. N and P may be used in the prevention, CC diagnosis and treatment of diseases associated with inappropriate P CC expression. For example, N and P may be used to treat disorders CC associated with decreased expression by rectifying mutations or deletions CC in a patient's genome that affect the activity of P by expressing CC inactive proteins or to supplement the patients own production of P CC Additionally, N may be used to produce the colon cancer-associated Ps, CC by inserting the nucleic acids into a host cell and culturing the cell CC begress the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 c and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 3428; 9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
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14-JUL-2000;
19-SEP-2000;
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                                                 and polynucleotides (II). (I) and (II) are useful for treating inflammatory conditions such as arthritis, nephritis, Crohn's disease, ischaemia-reperfusion injury, shock, sepsis, immune responses, and is involved in increasing haematopoiesis, stem cell survival, bone growth and remodeling. (I), (II) and modulators of (II) are useful for prophylaxis or treatment of one or more cancers (II) is also useful for creating transgenic animals useful for studying the in vivo activities of the polypeptide as well as for studying modulators of the polypeptides. (I) induces the proliferation of neural cells and regeneration of nerve and brain tissue and is useful for the treatment of central and peripheral nervous system diseases and neuropathies, such as Alzheimer's, Parkinson's disease, Huntington's disease, and amyotrophic lateral sclerosis. In addition, (I) is involved in chemotactic or chemokinetic activity, regulation of haematopoiesis and is useful for treating myeloid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel polypeptides and nucleic acids obtained from cDNA libraries prepared from various human tissues, for diagnosis and treatment cancer, neurological, inflammatory, and autoimmune disorders -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to novel isolated human secreted polypeptides (I)
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                            lymphoid cell disorders, platelet disorders such as thrombocytopenia
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DB; AAU28032.
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                                                                                                                                                 Human protein; hydrophobic domain; nutritional source; haematopoiesis; cytokine production; cell proliferation; cell differentiation; immune deficiency; infectious disease; autoimmune disorder; asthma; multiple sclerosis; systemic lupus erythematosus; rheumatoid arthritis; allergic reaction; osteoporosis; osteoarthritis; periodontal disease; nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; liver fibrosis; lung fibrosis; reperfusion injury; systemic cytokine damage; tissue differentiation; contraceptive; stroke; coagulation disorder; myocardial infarction; inflammatory condition;
                                                                                                                     coagulation disorder; myocardial infarction; septic shock; sepsis; ischaemia; reperfusion
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nephritis; therapy;

SS

injury; arthritis;

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Best Local Si
Query Match:
DB:
                                                                                                                                                       Percent Similarity:
                                                                                                                                                                                                CC infections. The protein is also used for treating autoimmune delicitions and to treat the first of the first of the first of all so used for treating autoimmune disorders conditions. The protein is also used for treating autoimmune disorders conditions such as asthma, and in immune suppression after organ conditions such as asthma, and in immune suppression after organ conditions such as asthma, and in immune suppression after organ conditions such as asthma, and in immune suppression after organ consequently in the treatment of myeloid or lymphoid cell conditions. The protein is useful in regulation of haematopoiesis conditions for the protein is also used in the treatment of osteoporosis consequently and in the treatment of periodontal disease and other contract processes. The protein is used in the treatment of nervous system disorders such as Alzheimer's disease, Parkinson's disease, and conditions resulting from systemic cytokine damage. They are useful for protection or regeneration and contraceptives since they exhibit activin or inhibin related conditions and as a fertility inducing therapeutic. They are used for treating various coagulation disorders and intreatment and prevention of conditions resulting from coagulation activities e.g. myocardial conditions resulting from coagulation activities e.g. myocardial conditions such as septic shock, sepsis, ischaemia correpert tumours.
                                                                                                                                                       Sequence 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence encodes a human protein of the invention, which has hydrophobic domains. The DNA sequences can be used as a probe or as a genetic marker. The protein can also be used as a marker, and to identify potential genetic disorders. The DNA and protein can also be used as nutritional sources or supplements. The protein exhibits cytokine, cell proliferation, cell differentiation activities and induces production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                other cytokines in certain cell populations. The protein also exhibits immune stimulating or immune suppressing activity. It can be used in the treatment of various immune deficiencies and disorders, and to treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human proteins having hydrophobic domains useful for treating osteoporosis, Alzheimer's disease, Parkinson's disease, asthma, multiple sclerosis, rheumatoid arthritis, cancer, anaemia, and stroke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; Page 217-218; 351pp; English.
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09-SEP-1998;
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07-AUG-1998;
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98JP-0224105.
98JP-0238116.
98JP-0254736.
98JP-0275505.
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US-09-801-115B-2 (1-99) x AAA15919 (1-297)

This sequence encodes a human protein of the invention, which has hydrophobic domains. The DNA sequences can be used as a probe or

be used as

a marker,

or as a to identify

Claim

4; Page 228; 351pp; English

multiple sclerosis,

rheumatoid arthritis, cancer,

anaemia,

stroke

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RESULT 13
AAA15929
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07-AUG-1998;
25-AUG-1998;
09-SEP-1998;
29-SEP-1998;
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                                                                                              Novel human proteins having hydrophobic domains useful for treating osteoporosis, Alzheimer's disease, Parkinson's disease, asthma,
                                                                                                                                                                                                                  (SAGA )
(PROT-)
                                                                                                                                             P-PSDB;
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                                                                                                                                                                                       Kato S,
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PROTEGENE INC.
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98JP-0224105.
98JP-0238116.
98JP-0254736.
98JP-0275505.
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RESULT 14
AAX97826
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Best Local Similarity:
Query Match:
DB:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC potential genetic disorders. The DNA and protein can also be used as cultitional sources or supplements. The protein exhibits cytokine, cell contercytokines in certain cell populations. The protein also exhibits comber cytokines in certain cell populations. The protein also exhibits content of various immune deficiencies and disorders, and to treat confectious diseases caused by viral, bacterial, fungal or other cell infections. The protein is also used for treating autoimmune disorders such as multiple sclerosis, systemic lupus erythematosus, and rheumatoid conditions such as asthma, and in immune suppression after organ conditions such as asthma, and in immune suppression after organ conditions such as asthma, and in immune suppression after organ transplantation. The protein is useful in regulation of haematopoiesis and consequently in the treatment of myeloid or lymphoid cell cell and consequently in the treatment of myeloid or lymphoid cell corosteoarthritis and in the treatment of proteodontal disease and other tooth repair processes. The protein is used in the treatment of osteoporosis or system disorders such as Alzheimer's disease, Parkinson's disease, and confictions resulting from systemic cytokine damage. They are used for treatment of lung or liver fibrosis, reperfusion injury in various system disorders and nother they exhibit activin or inhibin related activities and as a fertility inducing therapeutic. They are used for treating various coagulation disorders and in treatment and prevention of infarction or stroke. They also acts as receptors, receptor ligands or treat tunours.

Cc inhibitors or agonists of receptor/ligand interactions. They are used to crevent tunours.
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Human secreted protein encoding cDNA #14
                                        23-SEP-1999
                                                                                                             AAX97826 standard; cDNA; 500
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                                                                                                                                                                                                                                                                                                                                                                                                               174
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                                                                                                                                                                                                                                                                                                                                                                                                                              ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
                                                                                                                                                                                                                                                                            AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                                                                                                                                                                                                                       ValSerValLeuAlaLeuIleProGluThrThrThrLeuThrValGlyGlyGlyValPhe
                                                                                                                                                                                     PheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu 99
                                                                                                                                                                                                                                                          GCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTG
                                                                                                                                                                                                                                                                                                                                  GTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTT
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                                      (first entry)
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AAA38007
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13-NOV-1997;
17-DEC-1997;
09-FEB-1998;
13-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX97813-X97906 represent extended cDNA's which encode novel human secreted proteins (see AAY36129-Y36222) and which have cytostatic, thrombotic and osteopathic activity. The extended cDNAs can be used to express secreted proteins or parts of them or to obtain antibodies capable of binding to the secreted proteins. They may also be used in diagnostic, forensic, gene therapy and chromosome mapping procedures. Uses also include design of expression vectors and secretion vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 500 BP; 135 A; 102 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 28; Page 173-174; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Extended cDNAs encoding secreted proteins
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                                                                                                                                                                                                                               CGTATCTGTGTTGGCACTGATACCAGAAAACAACATTGACAGTTGGTGGAGGGGTGTT
                                                                              uPheAsnProSerGlyProTyrGlnLysLysProValHisGluLysLysGluValLeu
                                                                                                                                                      TGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCT
                                                                                                                                                                          eAlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuILeTyrArgLysLeuLe
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97US-0066677.
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98US-0074121.
98US-0081563.
98US-0096116.
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Indels:
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AAA38007 standard; cDNA; 459

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chemotaxic factor useful for treatment and diagnosis of immunocyte disorders - has immunocyte chemotaxic stimulating factor \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 459 BP; 123 A; 99 C;
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radiotherapy; chemotherapy; human; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
                       IleIleAsnSerLeuValThrThrValPheMetLeuIleValSerValLeuAlaLeuIle 47
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ATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATA
                                                                                                                                                                                                                                                                                                            GTGAAGATGCTGCGGCTGGCACTAACTGTGACATCTATGACCTTTTTTATCATCGCACAA 120
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CysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeuPheAsnProSerGlyProTyr
                                                                                                 ProGluThrThrTeuThrValGlyGlyGlyValPheAlaLeuValThrAlaValCys
                                          TGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTAC
  456
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Search completed: June 25, 2003, 12:51:07 Job time: 217 secs

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Result
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Fgapop 6.0 , Fgapext
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Listing first 45 summaries
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/B_COMB.seq:*
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US-09-385-982-92
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US-09-116-498-11
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US-09-103-840A-2
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Sequence 11, Appl
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Sequence 62, Appl	37,	37,	Sequence 37, Appl	Sequence 1, Appli	Sequence 9, Appli	Sequence 12, Appl	Sequence 12, Appl	Sequence 12, Appl	Sequence 12, Appl	Sequence 9, Appli	Sequence 14, Appl	Sequence 14, Appl	Patent No. 5320941	Sequence 213, App	Sequence 2, Appli	Sequence 319, App	Sequence 167, App	Sequence 108, App	Sequence 376, App	Sequence 1, Appli	Sequence 2, Appli	Sequence 1, Appli	Sequence 1, Appli	6	8	8,	176	39,	70,	Sequence 14, Appl	1,

#### ALIGNMENTS

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Sequence 98, Application Patent No. 6420526 EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/038,621
EARLIER FILING DATE: 1997-03-07 CURRENT APPLICATION NUMBER: US/ CURRENT FILING DATE: 1998-09-08 EARLIER EARLIER EARLIER EARLIER FILING DATE: 1997-03-07 EARLIER APPLICATION NUMBER: 60/ EARLIER FILING DATE: 1998-03-06 FILE REFERENCE: PZ002P1 TITLE OF INVENTION: EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER EARLIER APPLICATION NUMBER: 60/040,162 EARLIER APPLICATION NUMBER: PCT/US98/04493 INFORMATION: APPLICATION NUMBER: 60/047,597
FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,502
FILING DATE: 1997-05-23 FILING DATE: 1997-03-07
APPLICATION NUMBER: 60/040,163
FILING DATE: 1997-03-07
APPLICATION NUMBER: 60/047,600 APPLICATION NUMBER: 60/040,626
FILLING DATE: 1997-03-07
APPLICATION NUMBER: 60/040,334
FILLING DATE: 1997-03-07 APPLICATION NUMBER: 60/040,333 APPLICATION NUMBER: 60/047,615 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/040,336 FILING DATE: 1997-03-07 APPLICATION NUMBER: Rosen et 186 Human Secreted proteins 60/047,633 US/09/149,476

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ER APPLICATION NUMBER: 60/043,670
ER FILING DATE: 1997-04-11
ER APPLICATION NUMBER: 60/056,632
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ER APPLICATION NUMBER: 60/047,632
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                           EARLIER APPLICATION NUMBER: PCT/US98/04493
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EARLIER APPLICATION NUMBER: 60/040,333
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TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P1
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ER FILING DATE: 1997-08-22
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ER APPLICATION NUMBER: 60/056,908
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ER APPLICATION NUMBER: 60/048,964
ER FILING DATE: 1997-08-05
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ER APPLICATION NUMBER: 60/056,884
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/057,669
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ER APPLICATION NUMBER: 60/057,669
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ER APPLICATION NUMBER: 60/040,333
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APPLICATION NUMBER: 60/1
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ER FILING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,596
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RESULT 3
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VS-09-385-982-95
Sequence 95, Application US/09385982
Patent NO. 6262334
GENERAL IMPORMATION:
APPLICANT: ENDEGE, WILSON O., ET AL.
TITLE OF INVENTION: PRODUCTS: II
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ER APPLICATION NUMBER: 60/056,909
ER APPLICATION NUMBER: 60/056,909
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,875
EER APPLICATION NUMBER: 60/056,875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ER APPLICATION NUMBER: 60/048,964
ER FILING DATE: 1997-06-06
ER APPLICATION NUMBER: 60/057,650
ER APPLICATION NUMBER: 60/056,884
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/057,669
ER FILING DATE: 1997-09-05
ER APPLICATION NUMBER: 60/049,610
ER FILING DATE: 1997-06-13
ER APPLICATION NUMBER: 60/061,060
ER FILING DATE: 1997-06-13
ER APPLICATION NUMBER: 60/061,060
ER FILING DATE: 1997-10-02
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R APPLICATION NUMBER: 60/0:
R FILING DATE: 1997-08-22
R FILING DATE: 1997-08-22
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FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/
FILING DATE: 1997-08-22
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APPLICATION NUMBER: 60/043,670
PARTITUG DATE: 1997-04-11
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FILING DATE: 1997-08-22
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                                                                                                                                ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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AND GENE EXPRESSION

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80 356

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; FEATURE: misc_feature; LOCATION: (1)...(392); OTHER INFORMATION: n = US-09-385-982-95
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Best Local Similarity:
Query Match:
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CURRENT FILING DATE: 1999-08-30
EARLIER APPLICATION NUMBER: 09/328,111
EARLIER FILING DATE: 1999-06-08
EARLIER APPLICATION NUMBER: 60/117,393
EARLIER FILING DATE: 1999-01-27
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 544
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                                                                                       APPLICANT: Warren, Gregory W.
APPLICANT: Chen, Marcha
APPLICANT: Chen, Jeng S.
TITLE OF INVENTION: NOVEL INSECTICIDAL TOXINS FROM PHOTORHABDUS LUMINESCENS
TITLE OF INVENTION: AND NUCLEIC ACID SEQUENCES CODING THEREFOR
FILE REFERENCE: CGC1963/A
CURRENT APPLICATION NUMBER: US/09/251,645
CURRENT FILING DATE: 1999-02-17
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 11
LENGTH: 37948
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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11, Application US/09251645 Patent No. 6281413
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LENGTH: 392
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                                     TYPE: DNA ORGANISM: Photorhabdus luminescens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH:
NAME/KEY: CDS
LOCATION: (15171)..(18035)
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Indels:
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US-09-801-115B-2 (1-99) x US-09-251-645-11 (1-37948)
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Best Local Similarity:
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LOCATION: (31393)..(35838)
OTHER INFORMATION: Orf2
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LOCATION: (23768)...
OTHER INFORMATION:
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             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/116,498 FILING DATE: 16-UL-1998 CLASSIFICATION: UDknown>
                                                                                                                                                                   ZIP: 07601
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson, Esq.
STREET: 411 Hackensack Ave, Continental Plaza, 4th
                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: NOVEL ALTERNATIVE G-COUPLED RECEPTORS
ASSOCIATED WITH RETROVIRAL ENTRY INTO CELLS, METHODS
IDENTIFYING THE SAME, AND DIAGNOSTIC AND THERAPEUTIC
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
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RMATION: hph2
                                                                                                                                                                                                                                CITY: Hackensack
STATE: New Jersey
COUNTRY: USA
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Unutmaz, Derya
Ramani, Vineet N.K.
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                                                                                                        Version
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US-09-116-498-7; Sequence 7, Application US/09116498; Patent No. 6251582; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDENNESS: double
STRANDENNESS: double
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ORIGINAL SOURCE:
                         ZIP: 07601
COMPUTER READABLE FORM:
                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson,
                                                                                                                                                                        NUMBER OF SEQUENCES: 18
                                                                                                                                                                                                                                        TITLE
                                                                                                                                                                                                                                                                                                 APPLICANT: Littman, Dan R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: pigtail macaque SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 1049-1-009
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
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                                                                                       CITY: Hackensack
                                                         COUNTRY:
                                                                      STATE: New Jersey
                                                                                                                       STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCATATGGTCCCTGGTGGCCTTAATTTTCACCTTTTTTGTCCCTTTGTTG------
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      MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Jackson Esq., David A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 1083 base pairs
                                                                                                                                                                                            INVENTION: NOVEL ALTERNATIVE G-COUPLED RECEPTORS
ASSOCIATED WITH RETROVIRAL ENTRY INTO CELLS, METHODS OF
LDENTIFYING THE SAME, AND DIAGNOSTIC AND THERAPEUTIC USES
                                                                                                                                                                                                                                                  Unutmaz, Derya
Ramani, Vineet N.K.
                                                                                                                                                                                                                                                                                Deng, Hongkui
                                                                                                      Floor
                                                                                                                    411 Hackensack Ave, Continental Plaza, 4th
                                                         USA
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Indels:
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Matches:
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CITY: Alexandria
STATE: Virginia
                                 STREET:
                                                  ADDRESSEE:
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US-09-801-115B-2 (1-99) x US-09-116-498-7 (1-1083)
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                                                                                                                                                                                                                                          Sequence 2, Application US/08061092A Patent No. 5587458
                                                                                                                                                                                                                 GENERAL INFORMATION:
                                             NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                APPLICANT: KING, C R
APPLICANT: KASPRZYK, Philip G
APPLICANT: BIRD, ROBORT E
TITLE OF INVENTION: ANTI-CIBB-2 ANTIBODIES, COMBINATIONS
TITLE OF INVENTION: THEREOF, AND THERAPEUTIC AND DIAGNOSTIC USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 7:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/116,498 FILING DATE: 16-Jul-1998 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Jackson Esq., David A. REGISTRATION NUMBER: 26,742
George Mason Bldg., Washington & Prince Sts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 1049-1-009
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                    Burns, Doane, Swecker & Mathis
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COMPUTER READABLE FORM:

22313-1404

COUNTRY:

United States

MEDIUM TYPE:

Floppy disk

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Best Local Similarity:
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                                                                                                                                                                                                                                                                                   US-09-103-840A-2/c
                                                                                                                                                                                                                                      Sequence 2, Application US/09103840A Patent No. 6294328
                                                                                                                                                                                                                       GENERAL INFORMATION:
TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBERCULOSIS FILE REFERENCE: 24366-20007.00 CURRENT APPLICATION NUMBER: US/09/103,840A CURRENT FILING DATE: 1998-06-24 NUMBER OF SEQ ID NOS: 2 SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (703) 836-6620
TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 2:
ENCRY: CHARACTERISTICS:
                                                                                                                                         APPLICANT: FLEISCHMAN, ROBERT D.
APPLICANT: WHITE, Owen R.
APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Meuth, Donna M
REGISTRATION NUMBER: 36,
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 01
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Indels:
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Percent Similarity:
Best Local Similarity:
Query Match:
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; OTHER INFORMATION: "n" bases at various positions throughout the sequence
; OTHER INFORMATION: represent a, t, c or 9
US-09-103-840A-2
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US-09-801-115B-2 (1-99) x US-09-103-840A-1 (1-4411529)
                                                                   Query Match:
                                                                                     Percent Similarity:
Best Local Similarity:
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                                                                                                                                    Score
                                                                                                                                                           Pred. No.:
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                                                                                                                                                                                                                                                                                                                                  SEQ ID NO
                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: WHITE, Owen R.
APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBERCULOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: FLEISCHMAN, Robert D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Mycobacterium tuberculosis
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                                                                                                                                                                                                                                            ORGANISM: Mycobacterium tuberculosis OTHER INFORMATION: H37RV
                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                         ENGTH: 4411529
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Best Local Similarity:
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US-09-356-952-12
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Sequence 1,
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APPLICANT: Borlack-Sjodin, Ann
APPLICANT: Margarit, S. M.
APPLICANT: Bor-Sogi, Dafna
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SEQ ID NO 12
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CURRENT FILING DATE: 1999-07-19
EARLIER APPLICATION NUMBER: 60/093,631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EARLIER FILING DATE: 1998-07-21 NUMBER OF SEQ ID NOS: 14
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APPLICANT: Cole, Phillip
APPLICANT: Kuriyan, John
TITLE OF INVENTION: A CRYSTAL OF A RAS-SOS COMPLEX AND METHODS OF USE
TITLE OF INVENTION: THERBOF
FILE REFERENCE: 600-1-228N
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                                                                                                                                                                                                                                                43538 ATGAAAGTTAATGTGGATTTGTCCAAATTCGTCGATGGGCTTTACGCGCTATTACCTTAT 43479
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                                                                                                        85 GlyProTyrGlnLysLysPro-------ValHisGluLysLysGluValLeu 99
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                                                                                AATAACGAAATAATAAAACCATCAGTTAACGTTTCTACCAAAGCAGAACTTCTG 43365
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  Application US/09165396
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Matches:
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Best Local Similarity:
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APPLICANT: LUBKOWITZ, MARK A.
TITLE OF INVENTION: ISCLATED CANDIDA ALBICANS OLIGOPEPTIDE TRANSPORTER GENE
FILE REFERENCE: 372.6520p
CURRENT APPLICATION NUMBER: US/09/165,396
CURRENT FILING DATE: 1998-10-02
EARLIER APPLICATION NUMBER: PCT/US98/02332
EARLIER APPLICATION NUMBER: DCT/US98/02332
EARLIER FILING DATE: 1998-02-06
EARLIER FILING DATE: 1998-02-06
EARLIER APPLICATION NUMBER: 60/037,859
EARLIER APPLICATION NUMBER: 05/037,859
EARLIER FILING DATE: 1997-02-07
NUMBER OF SEQ ID NOS: 15
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                                                                     MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb sto COMEUTER: IBM AT-compatible, 80286 processor OPERATING SYSTEM: MS-DOS version 5.0 SOFTWARE: WordPerfect version 5.1 CURRENT APPLICATION DATA:
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LOCATION: (48
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                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: NO. 575:
TITLE OF INVENTION: Thereof
NUMBER OF SEQUENCES: 14
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APPLICANT: Hillen, Heinz
APPLICANT: Bialojan, Sie
                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                          STATE: I
                              FILING DATE:
               CLASSIFICATION:
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                                                      APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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1101 Connecticut Avenue
                                                                                                                                                                                                                          USA
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Matches:
Conservative:
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                                                    Alignment Scores:
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Best Local Similarity:
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                                     Pred. No.:
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Percent Similarity:
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FILING DATE: 19-JUL-1991
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 133 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09462844 Patent No. 6258563 GENERAL INFORMATION:
                                                                                                                                                              SEQ ID NO 1
                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/462,844
CURRENT FILING DATE: 2000-01-13
PRIOR APPLICATION NUMBER: PCT/US98/14786
PRIOR FILING DATE: 1998-07-15
PRIOR APPLICATION NUMBER: EP 97305286.3
PRIOR APPLICATION NUMBER: EP 97305344.0
                                                                                                                                                          PRIOR FILING .... NUMBER OF SEQ ID NOS: 11
NUMBER OF SEQ ID NOS: 11
FRANKSEQ for Windows Version 4.
                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Increasing Production of Proteins TITLE OF INVENTION: Gram-Positive Microorganisms FILE REFERENCE: GC385-US
                                                                                                                                                                                                              PRIOR FILING DATE:
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                                                                                                    LENGTH: 2211
TYPE: DNA
ORGANISM: Bacillus subtilis
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                                                                                                                                                                                                          UMBER: EP 97305344.0
1997-07-17
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              Length:
Matches:
Conservative:
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Indels:
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RESULT 14
US-08-772-270A-14/c
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Alignment Scores:
                                      ORGANISM:
US-08-772-270A-14
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                                                                                                                                                         TELEPHONE: (416) 364-7311
TELEFAX: (416) 361-1398
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 7721 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 14, Application US/08772270A Patent No. 6019984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 65
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION: NAME: Gravelle, Micheline REGISTRATION NUMBER: 40,2
                                                                             ORIGINAL SOURCE:
                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NOVEL BACTERIAL PREPARATIONS, METHOD FOR TITLE OF INVENTION: PRODUCING SAME, AND THEIR USE AS VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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APPLICANT: Rosendal, Soren
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CITY: Toronto
STATE: Ontario
                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: DOCUMENT OF CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/772,270A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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M5H 3Y2
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Best Local Similarity:
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NUMBER OF SEQUENCES: 142

CORRESPONDENCE ADDRESS:
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                                                                      TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                  COMPUTER: HP Vectra 486/33

OPERATING SYSTEM: MSDOS version 6.2

SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBERS
SEQUENCE CHARACTERISTICS:
LENGTH: 17710 base pairs
TYPE: nucleic acid
                                                                                                           REGISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/031,626 AND
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                             APPLICATION NUMBER: UPPLICATION NUMBER: UPPLICATION: 536
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                                                                                                                                                                                     Steffe, Eric K.
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Search completed: June 25, 2003, 13:33:51 Job time: 1226 secs
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DB:
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TOPOLOGY: lir
                                                                         17406 GCAGGTGGTGCTGTCGGGTCAGTGGCGGGGGGATATTGTTCAGTCCACTGGGACCATTT 17465
                                                                                                                                             17346 CGGGGACGGATCTCCGGTGCGCAGTTCATCAGGAATCTTGCCGTGGCATCTTCCTGTGTG 17405
                                                                                                                                                                                                                 17286 CTGGTGACATCTCTGGCACTGGTAGCTGTCACAACCGGCCCTGACATGATCAAAATGTTG 17345
                                                                                                                                                                                                                                                                                   17226 CTTCAGAAGGGGACCGGTGCAGGAAATATCAGTGCCCTGAACAAAGTGATGAAGGGGTCG 17285
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                                                                                                      AlaAspGlyAlaLeu-----IleTyrArgLysLeuLeuPheAsnProSerGlyProTyr 87
                                                                                                                                                                            ThrLeuThrValGlyGlyGlyValPheAla-----LeuValThrAlaValCysCysLeu
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Command line parameters:

-MODEL-frame+p2n.model -DEV=xlp
-MODEL-frame+p2n.model -DEV=xlp
-Q-/cgn2_1/USPTO_Spool_VUS9801115/runat_20062003_141105_13685/app_query.fasta_1.263
-DB=Published_Applications_NA -OFMT-fastap -SUFFIX=rnpb -MINMATCH=0.1
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MX=100
-THR_MIN=0 -ALICK=15 -MODE-LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXLEN=2000000000 -USER-US09801115_@CGN_1 180_@runat_20062003_141105_13685
-NCPU=6 -ICPU=3 -NO_MAAP -LARGEDUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100
-LONGLOG -DEV_TIMEDUT=120 -WARN_TIMEDUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOPP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                          Score
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Match
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:/cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
:/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
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:/cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
:/cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
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US-09-822-846-397
US-09-809-391-98
US-09-809-391-252
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Sequence 6534, Ap
Sequence 397, App
Sequence 98, Appl
Sequence 252, App
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		1927, 1927, 19489 189, 189, 189, 189, 189, 189, 189, 18

### ALIGNMENTS

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RESULT 1
US-09-918-995-6534
                                   Percent Similarity:
Best Local Similarity:
                                                                      Score:
                                                                                      Pred. No.:
                 Query Match:
                                                                                                        Alignment Scores:
                                                                                                                                          US-09-918-995-6534
                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6534, Application US/09918995 Publication No. US20030073623A1 GENERAL INFORMATION:
                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 6534
                                                                                                                                                                                                                                                                                             APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
                                                                                                                                                                        LENGTH: 422
TYPE: DNA
                                                                                                                                                        ORGANISM: Homo sapiens
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                                   Conservative: Mismatches:
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                                          US-09-801-115B-2 (1-99) x US-09-822-846-397 (1-439)
                                                                                       Query Match:
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                                                                                                                                                                                                   ; SOFTWARE: PatentIn Ver. 2; SEQ ID NO 397; LENGTH: 439; TYPE: DNA; ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                       APPLICANT: Gulukota, Kamalakar
APPLICANT: Graham, James R.
APPLICANT: Graham, James R.
APPLICANT: Genetics Institute, Inc.
APPLICANT: Genetics Institute, Inc.
APPLICATION: POLYNUCLEOTIDES ENCODING NOVEL SECRETED PROTEINS
FILLE REFERENCE: GIN 6400
CURRENT APPLICATION NUMBER: US/09/822,846
CURRENT FILING DATE: 2001-03-29
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR APPLICATION DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 629
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MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
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Wong, Gordon G.
Clark, Hilary
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Steininger II, Robert
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Sequence 252, Application US/09809391
Publication No. US20030049618A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
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RESULT 4
US-09-809-391-252
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US-09-809-391-98
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Publication No. US20030049618A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 98
LENGTH: 504
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/809,391 CURRENT FILING DATE: 2001-03-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo
                                                         131 ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
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                               ; TYPE: DNA
; ORGANISM: HOMO
US-09-801-115-1
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Alignment Scores:
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APPLICANT: Song, Q.
APPLICANT: bil, C.
APPLICANT: Huang, J.
APPLICANT: Tang, J.
APPLICANT: Tang, J.
APPLICANT: Chen, G.
TITLE OF INVENTION: CHEMOKINE LIKE FACTOR(CKLF) WITH CHEMOTACTIC
TITLE OF INVENTION: STIMULATING ACTIVITIES
TITLE OF INVENTION: US/09/801,115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 252
LENGTH: 506
                                                                                                              FILE REFERENCE: 10776-003-999
FILE REFERENCE: 10776-003-999
CURRENT FILING DATE: 2001-03-07
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: PCT/CN00/00026
PRIOR FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 99107284.7
PRIOR FILING DATE: 1999-05-14
                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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                                                                                            SEQ ID NO 1
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CURRENT FILING DATE: 2001-03-16
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                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: n equals a,t,g, US-10-106-698-1927
                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/106,698
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: PCT/US00/26524
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US 60/157,137
PRIOR APPLICATION NUMBER: US 60/163,280
PRIOR FILING DATE: 1999-09-29
PRIOR APPLICATION NUMBER: US 60/163,280
PRIOR FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 8564
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LENGTH: 558
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TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypept
FILE REFERENCE: PA005P1
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                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature LOCATION: (514)..(514)
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 21 ValLysMetLeuArgLeuAspIleIleAsnSerLeuValThrThrValPheMetLeuIle 40
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US-09-801-115-3; Sequence 3, App.; Patent No. US2
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TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 2041-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: US/09/235,076
PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 19489
LENGTH: 512
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NAME/KEY: misc_feature

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; OTHER INFORMATION: n = A,T,C

US-09-918-995-19489
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          ce 3, Application US/09801115
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                                                                                          TTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAAAANAAGAAGTTTTG
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Sequence 189, Application US/09978295A Patent No. US20020156006A1 GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: CHEMOKINE LIKE FACTOR(CKLF) WITH CHEMOTACTIC TITLE OF INVENTION: STIMULATING ACTIVITIES FILE REFERENCE: 10776-03-999 CURRENT APPLICATION NUMBER: US/09/801,115 CURRENT FILING DATE: 2001-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1999-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/CN00/00026
PRIOR FILING DATE: 2000-02-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 GCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTTATTTTTCATA 180
                                                                                                                                  88 GlnLysLysProValHisGluLysLysGluValLeu 99
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                                                                                                                    CAGAAAAAGCCTGTGCATGAAAAAAAAAGAAGTTTTG
                                                                                                                                                                                                     CysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeuPheAsnProSerGlyProTyr
                                                                                                                                                                                                                                                                        ProGluThrThrThrLeuThrValGlyGlyGlyValPheAlaLeuValThrAlaValCys
                                                                                                                                                                                     TGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTAC
                                                                                                                                                                                                                                                     CCAGAAACCACAACATTGACAGTTGGTGGAGGGGGTGTTTGCACTTGTGACAGCAGTATGC
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Han, W.
Zhang, Y.
Song, Q.
Di, C.
Huang, J.
Tang, J.
Chen, G.
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Conservative:
Mismatches:
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FILE REFERENCE: P2650PICIL
CURRENT APPLICATION NUMBER: US/09/978,295A
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: 09/062250
PRIOR FILING DATE: 2097-01-17
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-11-33
PRIOR APPLICATION NUMBER: 60/06331
PRIOR FILING DATE: 1997-11-33
PRIOR APPLICATION NUMBER: 60/06364
PRIOR FILING DATE: 1997-11-31
PRIOR APPLICATION NUMBER: 60/077450
PRIOR FILING DATE: 1997-11-21
PRIOR APPLICATION NUMBER: 60/077632
PRIOR FILING DATE: 1998-03-10
PRIOR FILING DATE: 1998-03-11
PRIOR APPLICATION NUMBER: 60/077641
PRIOR APPLICATION NUMBER: 60/077641
PRIOR APPLICATION NUMBER: 60/077641
PRIOR APPLICATION NUMBER: 60/077641
PRIOR APPLICATION NUMBER: 60/077649
PRIOR APPLICATION NUMBER: 60/077649
PRIOR FILING DATE: 1998-03-11
PRIOR APPLICATION NUMBER: 60/07791
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NPPLICANT: Williams, P. Mickey
NPPLICANT: Wood, William I.
NPLICANT: Wood, William I.
NPLICANT: No. Secreted and Transmembrane Polypeptides and Nucleic
NITLE OF INVENTION: Acids Encoding the Same
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FILING DATE: 1998-03-27
APPLICATION NUMBER: 60/079728
FILING DATE: 1998-03-27
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APPLICATION NUMBER: 60/078939
FILING DATE: 1998-03-20
APPLICATION NUMBER: 60/079294
FILING DATE: 1998-03-25
FILING DATE: 1998-03-25
                                                                                APPLICATION NUMBER: 60/
FILING DATE: 1998-03-27
APPLICATION NUMBER: 60/
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APPLICATION NUMBER: 60/
FILING DATE: 1998-03-27
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Grimaldi, J. Christopher
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Stewart, Timothy A.
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Kuo, Sophia S.
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Kljavin, Ivar J.
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PRIOR APPLICATION NUMBER: 60/080333
PRIOR FILING DATE: 1998-04-01 PRIOR FILING DATE: 1998-04-15
PRIOR APPLICATION NUMBER: 60/081952
PRIOR FILING DATE: 1998-04-15
PRIOR APPLICATION NUMBER: 60/081838
PRIOR FILING DATE: 1998-04-15 OR APPLICATION NUMBER: 60/081195
OR FILING DATE: 1998-04-08
OR APPLICATION NUMBER: 60/081203
OR ETLING DATE: 1998-04-09
OR APPLICATION NUMBER: 60/081229
OR FILING DATE: 1998-04-09
OR FILING DATE: 1998-04-15
OR FILING DATE: 1998-04-15
OR APPLICATION NUMBER: 60/081817
OR FILING DATE: 1998-04-15
OR FILING DATE: 1998-04-15
OR FILING DATE: 1998-04-15 DR APPLICATION NUMBER: 60/080194
DR FILING DATE: 1998-03-31
DR APPLICATION NUMBER: 60/080327
DR FILING DATE: 1998-04-01
DR APPLICATION NUMBER: 60/080328
DR FILING DATE: 1998-04-01
DR FILING DATE: 1998-04-01 DR FILING DATE: 1998-04-22

DR APPLICATION NUMBER: 60/082797

DR FILING DATE: 1998-04-22

DR APPLICATION NUMBER: 60/082796

DR FILING DATE: 1998-04-23

DR APPLICATION NUMBER: 60/083336

DR APPLICATION NUMBER: 60/083336 OR FILING DATE: 1998-04-29

R APPLICATION NUMBER: 60/083499

OR FILING DATE: 1998-04-29

OR APPLICATION NUMBER: 60/083545

OR FILING DATE: 1998-04-29

OR APPLICATION NUMBER: 60/083554 FILING DATE: 1998-04-08
APPLICATION NUMBER: 60/081071
FILING DATE: 1998-04-08 APPLICATION NUMBER: 60/081049 APPLICATION NUMBER: 60/081070 FILING DATE: 1998-04-08 FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: 60/ FILING DATE: 1998-03-31 FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: 60/079786 FILING DATE: 1998-03-27 APPLICATION NUMBER: 60/082804 FILING DATE: 1998-04-22 APPLICATION NUMBER: 60/082700 APPLICATION NUMBER: 60/082569
FILING DATE: 1998-04-21
APPLICATION NUMBER: 60/082704
FILING DATE: 1998-04-22 APPLICATION NUMBER: 60/082568 FILING DATE: 1998-04-21 FILING DATE: 1998-04-01 FILING DATE: 1998-03-APPLICATION NUMBER: FILING DATE: 1998-0 APPLICATION NUMBER: APPLICATION NUMBER: 60/083495 FILING DATE: 1998-04-29 APPLICATION NUMBER: 60/083496 FILING DATE: 1998-04-27 APPLICATION NUMBER: 60/083322 FILING DATE: 1998-04-28 APPLICATION NUMBER: 60/081819 APPLICATION NUMBER: 60/080334 APPLICATION NUMBER: 60/083392 FILING DATE: 1998-04-29 1998-03-31 1998-03-1998-03-30 60/080105 60/080165 60/080107 60/079923 60/079920

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                                                                                                                                                                                                                                                                                                               Pred. No.:
                                                                                                                                                                                                                                                                                 Percent Similarity:
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R FILING DATE: 1998-05-06

DR APPLICATION NUMBER: 60/084441

DR FILING DATE: 1998-05-06

DR APPLICATION NUMBER: 60/084637

DR FILING DATE: 1998-05-07
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OR FILING DATE: 1998-04-29
OR APPLICATION NUMBER: 60/083742
OR FILING DATE: 1998-04-30
OR APPLICATION NUMBER: 60/084366
OR FILING DATE: 1998-05-05
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FILING DATE: 1998-05-15
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FILING DATE: 1998-05-07
APPLICATION NUMBER: 60/084627
FILING DATE: 1998-05-07
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FILING DATE: 1998-05-07
APPLICATION NUMBER: 60/084640
FILING DATE: 1998-05-07
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APPLICATION NUMBER: 60/
FILING DATE: 1998-04-29
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                 251 GCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTTATTTTTCATA 310
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                                                                           GTGAAGATGCTGCGGCTGGCACTAACTGTGACATCTATGACCTTTTTTATCATCGCACAA
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1998-05-15
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1998-05-15
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Matches:
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US-09-978-697-189
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PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/066364
PRIOR FILING DATE: 1997-11-21
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PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/064249
PRIOR FILING DATE: 1997-11-03
                                                                                                                                                                                                                                                                     APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2630P1C27
                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/978,697
CURRENT FILING DATE: 2001-10-16
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PRIOR FILING DATE: 2001-07-30
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APPLICANT: Baker Kevin p
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Williams, P. Mickey
Wood, William I.
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Stewart, Timothy A.
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Hillan, Kenneth J
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Grimaldi, J. Christopher
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Goddard, Audrey
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Filvaroff,
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FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
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                                                       Kljavin, Ivar J.
                                                                                        Gurney, Austin L.
Hillan, Kenneth J
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Grimaldi, J. Christopher
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                                         Napier, Mary A.
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PRIOR APPLICATION NUMBER: 60/066364
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APPLICATION NUMBER: 60/079728
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Stewart, Timothy A.
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DR FILING DATE: 1998-04-15
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                            DR FILING DATE: 1998-03-20
DR APPLICATION NUMBER: 60/079294
DR FILING DATE: 1998-03-25
DR APPLICATION NUMBER: 60/079656
DR FILING DATE: 1998-03-26
DR APPLICATION NUMBER: 60/079664
                                                                                                                                                                                  OR FILING DATE: 1998-03-12
OR APPLICATION NUMBER: 60/078004
OR FILING DATE: 1998-03-13
OR APPLICATION NUMBER: 60/078886
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OR APPLICATION NUMBER: 60/078936
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OR FILING DATE: 1997-11-03
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Shelton, David L.
Stewart, Timothy A.
Tumas, Daniel
Williams, P. Mickey
Wood, William I.
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Hillan, Kenneth J
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     OR APPLICATION NUMBER: 60/083322
OR FILING DATE: 1998-04-28
OR APPLICATION NUMBER: 60/083392
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FILING DATE: 1998-04-22
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APPLICATION NUMBER: 60/082804
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FILING DATE: 1998-04-15
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APPLICATION NUMBER: 60/081203
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APPLICATION NUMBER: 60/079663
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APPLICATION NUMBER: 60/079728
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FILING DATE: 1998-04-09
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FILING DATE: 1998-04-08
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Percent Similarity:
Best Local Similarity:
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Pred. No.:
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RAPPLICATION NUMBER: 60/083742

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FILING DATE: 1998-05-15
APPLICATION NUMBER: 60/085700
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APPLICATION NUMBER: 60/085323
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PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/064249
PRIOR FILING DATE: 1997-11-03
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
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CURRENT FILING DATE: 2001-10-15
PRIOR APPLICATION NUMBER: 09/918585
PRIOR FILING DATE: 2001-07-30
                                                                                                                                                                                                                                             APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Tra
TITLE OF INVENTION: Acids Encoding
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Shelton, David L.
Stewart, Timothy A.
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Grimaldi, J. Christopher
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OR APPLICATION NUMBER: 60/080194
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PRIOR APPLICATION NUMBER: 60/ FILING DATE: 1998-05-13 FILING DATE: 1998-0 APPLICATION NUMBER: FILING DATE: APPLICATION APPLICATION NUMBER: 60/ FILING DATE: 1998-05-15 FILING DATE: 1998-05-05 APPLICATION NUMBER: 60/0 FILING DATE: 1998-05-06 FILING DATE: 1998-04-29
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           Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
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                                                                      Gerber, Hanspeter
Gerritsen, Mary E.
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 Kljavin,
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          Sequence 189, Application US/09978191A
Publication No. US20030050239A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi
APPLICANT: Baker Kevin P.
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APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P263091C22
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                                                                                                                    GlnLysLysProValHisGluLysLysGluValLeu 99
                                                                                                                                                      GTGAAGATGCTGCGGCTGGCACTAACTGTGACATCTATGACCTTTTTTATCATCGCACAA 250
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Shelton, David L.
Stewart, Timothy A.
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Paoni, Nicholas F
Botstein, David
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CURRENT FILING DATE: 2001-10-15
PRIOR APPLICATION NUMBER: 09/918585
PRIOR FILING DATE: 2001-07-30
PRIOR FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: 60/062250
PRIOR APPLICATION NUMBER: 60/064249
PRIOR APPLICATION NUMBER: 60/064249
PRIOR FILING DATE: 1997-11-03
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/06564
PRIOR APPLICATION NUMBER: 60/077450
PRIOR APPLICATION NUMBER: 60/077452
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APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
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OR FILING DATE: 1998-03-11
OR APPLICATION NUMBER: 60/077641
OR APPLICATION NUMBER: 60/077649
OR APPLICATION NUMBER: 60/0777649
OR APPLICATION NUMBER: 60/077791
OR APPLICATION NUMBER: 60/07791
OR APPLICATION NUMBER: 60/078886
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OR FILING DATE: 1998-03-20
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Stewart, Timothy A.
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Godowski, Paul J.
Grimaldi, J. Christopher
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OR APPLICATION NUMBER: 60/0
OR AFFILING DATE: 1998-04-21
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OR AFFILING DATE: 1998-04-21
                                                                                                                                                                                                                                                                                                                                   FILING DATE: 1998-04-22
APPLICATION NUMBER: 60/
FILING DATE: 1998-04-23
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APPLICATION NUMBER: 60/081952
FILING DATE: 1998-04-15
APPLICATION NUMBER: 60/081838
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FILING DATE: 1998-04
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APPLICATION NUMBER: 60/081229
FILING DATE: 1998-04-09
APPLICATION NUMBER: 60/081955
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APPLICATION NUMBER: 60/081070
FILING DATE: 1998-04-08
                                     APPLICATION
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FILING DATE:
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APPLICATION NUMBER: 60/081203
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NUMBER: 60/1: 1998-04-29 NUMBER: 60/ : 1998-04-29 NUMBER:

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60/082700 60/082804

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APPLICATION NUMBER: 60/083500 FILING DATE: 1998-04-29 APPLICATION NUMBER: 60/083742 FILING DATE: 1998-04-30

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FILING DATE: 1998-05-05
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	88 GlnLysLysProValHisGluLysLysGluValLeu 99	8	Qy
550	491 TGTCTTGCCGACGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTAC 550	49	Дb
87	68 CysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeuPheAsnProSerGlyProTyr 87	6	VQ.
490	431 CCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGC 490	43	DЬ
67	48 ProGluThrThrLeuThrValGlyGlyGlyValPheAlaLeuValThrAlaValCys 67	4	ρ
430	371 ATTATCAACTGATAACAACAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATA 430	37	В
47	28 IleIleAsnSerLeuValThrThrValPheMetLeuIleValSerValLeuAlaLeuIle 47	2	Qy
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Scoring table: Sequence: Title: Perfect score: Run on: OM nucleic - nucleic search, using sw model June 25, 2003, 06:26:50 ; Search time 1523 Seconds (without alignments) 10204.135 Million cell updates/sec IDENTITY\_NUC
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Searched:

2054640 segs, 14551402878 residues

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database GenEmbl:\* gb\_pr:\* gb\_pat:\*
gb\_ph:\*
gb\_pl:\* gb\_ba:\* gb\_htg:\* gb\_ov:\* gb\_sts:\*
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Pred. No. is the number of results predicted by chance to have

em\_htgo\_other:\* em\_htgo\_hum:\*
em\_htgo\_mus:\*

em\_htg\_vrt:\*
em\_sy:\*

em\_htg\_mam: \*

em\_htg\_pln:\* em\_htg\_rod:\* em\_htg\_other: \* em\_htg\_inv:\*

em\_htg\_mus:\*

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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## ALIGNMENTS

RESULT 1
AF096895
LOCUS
DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE
ORGANISM REFERENCE AUTHORS

em\_un:\* em\_vi:\*

530 bp mRNA linear PRI 18-JUL-2.
Homo sapiens chemokine-like factor 1 (CKLF1) mRNA, complete cds.
AF096895
AF096895.2 GI:9989691
FLI\_CDNA. Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 530)
Han, W., Lou, Y., Tang, J., Zhang, Y., Chen, Y., Li, Y., Gu, W., Huang, J.,
Gui, L., Tang, Y., Li, F., Song, Q., Di, C., Wang, L., Shi, Q., Sun, R.,
Xia, D., Rui, M., Tang, J. and Ma, D. Homo sapiens. PRI 18-JUL-2001

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Han,W.L., Li,Y., Zhang,Y.M., Di,C.H., Song,Q.S. and Ma,D.L.
Direct Submission
Submitted (03-0CT-1998) Immunology, Beijing Medical University,
Xue Yuan Rd., Beijing 100083, China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Science Center, xueyuan Road 38, Beij
Sequence update by submitter
On Sep 7, 2000 this sequence version
Location/Qualifiers
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Biochem. J. 357 (Pt 1), 127-135 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Molecular cloning and characterization of chemokine-like factor 1 (CKLF1), a novel human cytokine with unique structure and potential
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                            TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
                                                                                          TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
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ALIPETTTLTVGGGVEALVTAVCCLADGALIYRKLLENPSGPYQKKPVHEKKEVL"
1 108 c 137 g 129 t
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expression inhibited by IL-10"
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/protein_id="AAF06722.1"
/db_xref="GI:6288734"
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/db_xref="taxon:9606"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; H
1 (bases 1 to 515)
dumas milne Edwards, J.B., Bougueleret, L. and Jobert, S.
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/LIPETTTITVGGGVFALVTAVCCLADGALIVRKLLFNPSGPYQKKPVHEKKEVL"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
Submitted (17-MAR-1999) Immunology, Beijing Medical University, Xue
Yuan Road, Beijing 100083, China
2 (bases 1 to 689)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             alternatively spliced AF135380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AF135380 689 bp mRNA linear Homo sapiens chemokine-like factor 2 (CKLF2) mRNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (07-SEP-2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Han, W.L., Gu, W., Li, Y., Zhang, Y., Song, Q., Di, C. and Ma, D.
                                                                                                                                                                                                                                             Similarity
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                                                                                              ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG
                                                                                                                                                                             GTGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
                                                                                                                               ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGGAAAGTGCTGCTG
TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTG----
                                 CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT
                                                CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT
                                                                                                                                                            GGGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
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                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                      /translation-"MDNVQPKIKHRPEGESVKGHVKMLRLAITVTSMTEFIIAQAPEP
/TVITGEFEVTVILFEILLYVLBLDRLMKWLFWFLLDIINSLVTTVFMLIVSVLALIPE
TTTLTVGGGVEALVTAVCCLADGALYRKLLFNPSGPYQKKPVHEKKEVL"
1 140 c 161 g 193 t
                                                                                                                                                                                                                                                                                                                                                                   /product="chemokine-like factor 2"
/protein_id="AAF19599.1"
/db_xref="GI:6630854"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /gene="CKLF2"
/note="UCK"
                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="UCK-2; alternatively spliced"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"
/db_xref="taxon:9606"
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76.4%;
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Pred. No. 2.7e-75;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AF145216 593 bp mRNA linear Homo sapiens chemokine-like factor 4 (CKLF4) mRNA,
                                                                                                                                                                                                                                                                                          Submitted (23-APR-1999)
Yuan Road, Beijing 10008
2 (bases 1 to 593)
                                                                                                                                                                                                                                                                                                                                                         Ma, D.
                                                                                                                                                                             Sequence update by submitter
On Sep 7, 2000 this sequence version replaced gi:6625671
Location/Qualifiers
                                                                                                                                                                                                                            Submitted (07-SEP-2000) Immunology, Peking University Health Science Center, Xueyuan Road 38, Beijing 100083, China
                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                              Direct Submission
                                                                                                                                                                                                                                                                             Han, W.L
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                         (bases .1 to 593)
                                                                                                                                                                                                                                                                                                                                                                    W.L., Gu,W.F.,
                148. .510
/gene="CKLF4"
                                                                   /gene="CKLF4"
                                                                                                                /cell_line="U937"
                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                   /note="from PHA
     /note="UCK-4; alternatively spliced"
                                                    'note="UCK"
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                                                                                                                                                                                                                                                                                                             100083,
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                                                                                                                                                                                                                                                                                                                         Immunology,
                                                                                                    stimulated cells"
                                                                                                                                                                                                                                                                                                             China
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395 490

550

430 275 370 229

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SOURCE
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                                                                                               REFERENCE
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                                                                            AUTHORS
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Direct Submission
Submitted (12-MAR-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11
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                                                                                                                                                                                                                    Homo sapiens, clone MGC:10658 BC004380
                                                                           Strausberg, R.
                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                Homo sapiens
                                                                                                                                                                                     MGC
                                                                                                                                                                                                BC004380.1 GI:13325133
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                                                                                                                                                                 Homo sapiens.
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                                                                                           (bases 1 to 688)
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YIVITGFEVTVILFFILLYVLRLDRLMKWLFWPLLVFALVTAVCCLADGALIYRKLLF
NPSGYYQKKPVHEKKEVL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /product="chemokine-like factor 4"
/protein_id="AAF19350.1"
/db_xref="GI:6625672"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /codon_start=1
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79.8%;
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Pred. No. 3.4e-75;
0; Mismatches 55;
                                                                                                                                                                                                                      -----ATATTATCAACTCACTGGTAACAACAG
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Best Local Similarity
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                                             313 CATACTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCCTTTGCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hsiao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prabhu, Parvaneh Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy, George Yang, Scott Zuyderduyn, Marco Marra.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA Library Preparation: Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Cou DNA Sequencing by: Genome Sequence Centre, BC Cancer Agency, Vancouver, BC, Canada info@bcgsc.bc.ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-remail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGCTGGGTCTGCAGA 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGCCGAGAAGTAGGGGAGGCCGCTGCTCCGCCGCGGTTGCTATCGCTTCGCAGA
-GATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTTGGCACT
                                                                                                                                          ACAAGCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTTATTTTT 312
                                                                                                                                                                                                                                          CCACGTGAAGATGCTGCGGCTGGCACTAACTGTGACATCTATGACCTTTTTTATCATCGC
                                                                                                                                                                                                                                                                                               CCACGTGAAGATGCTGCGGCTG-----
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                                                                                                                                                                                                                                                                                                                                          CGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGG
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/protein_id="AAH04380,1"
/protein_id="AAH04380,1"
/db_xref="GI:13325134"
/translation="MONYOPKIKHRPECFSVKGHVKMLFLALTVTSMTFFIIAQAPEP
YIVITGFEVTVILFFILLYVLRLDRLMKWLFWFLLDIINSLVTTVFMLIVSVLALIPE
TTTLYYGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
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137. .595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="MGC:10658 IMAGE:3639550"
/tissue_type="Pancreas, adenocarcinoma"
/clone_lib="NHH_MGC_39"
/lab_host="DH10B-R"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No. 1.7e-72;
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61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (03-APR-1998) Pathology and Lab. and Foundation, Rochester, MN 55905, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 655)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zhang, J.S., Nelson, M., Wang, L. and Smith D.
                                                                                                                                     Similarity
                                                                          CCGAGCTGGGCGAGAAGTAGGGGAGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTTGCACTTGTGACAGCAGT
                          TCGCAGAACCTACTCAGGCAGCCAGCTGAGAAAAGTTGAGGGAAAGTGCTGCTGGGT 140
                                                         CCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCT 60
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TCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGGGT
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                                                                                                                                                                                                 171
                                                                                                                      Conservative
                                                                                                                                                                                               /translation="mdnvQpkikhrpfcfsvkGhvkmLrlaltvtsmfffiiaQapep
YIVITGfEVTYVLFffiltyVLRLDRLMKWLFWPLLDIINSLVTTVFMLIVSVLALIPE
TTTLTVGGGVFALVTAVCCLADGALTXKLLFNPSGPYQKKPVHEKKEVL"
137 c 153 g 194 t
                                                                                                                                                                                                                                                          /product="transmembrane proteolipid"
/protein_id="AAF21255.1"
/db_xref="GI:6648619"
                                                                                                                                                                                                                                                                                                                     sodium butyrate"
                                                                                                                                                                                                                                                                                                                                                   /gene="C32"
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132. .590
                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_line="AsPC-1"
                                                                                                                                                                                                                                                                                                      /codon_start=1
                                                                                                                                                                                                                                                                                                                                    /note="down regulated upon
                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="pancreas"
/note="adenocarcinoma"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
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75.6%;
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Pred. No. 2.1e-69;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 70
AX061665
                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                              178
                                                                                                                                                                                                                                                                                        (FR)
                                      /note="Von Heijne 630. .635 655. .669
                                                                                                  /codon_start=1
/protein_id="CAC25025.1"
/protein_id="CAC25025.1"
/db_xref="GI:1406790"
/translation="MONVQPKIKHRPECESVKGHVKMLRLALTVTSMTFFIIAQAPEP
YIVITGFEVTVILFFILLYVLRLDRLMKWLFWPLLDIINSLVTTVFMLIVSVLALIPE
YIVITGFEVTVLLFFILLYVLRLDRLMKKLLFNPSGFYQKKPVHEKKEVL"
                                                                                        140.
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/db_xref="taxon:9606"
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     Han, W.L
Direct
                                    Direct Submission
Submitted (17-MAR-1999) Immunology,
Yuan Road, Beijing 100083, China
                                                                                                                                                                                     alternatively spliced AF135381
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                                                                             1 (bases 1 to 434)
Han, W.L., Gu, W., Li, Y., Zhang, Y.,
                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                             Homo sapiens chemokine-like
                                                                                                                                               Homo sapiens.
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                              (bases 1 to 434)
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Pred. No. 1.6e-68;
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3 (CI
                                                      Beijing Medical University,
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G30204.1 GI:1593755
STS; STS sequence; I
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On Sep 7, 2000 this sequence version replaced gi:6630855.
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                                                                                                                                                                                                                                                                                                                                                                                                                               CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT
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                                                                                                                       TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
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/protein_id="AAF19600.1"
/db_xref="GI:6630856"
/translation="MUNYOPKIKHRPFCFSVKGHVKMLRLVFALVTAVCCLADGALIY
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/db_xref="taxon:9606"
/cell_line="U937"
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Primer B: GCCCTTATTTACCGGAAGCT
STS size: 77
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Myers
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Location/Qualifiers
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Mammalia; Eutheria;
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                                                               AGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (10-AUG-2001) DOE Joint Genome Institute, 2800 Mit Drive, Walnut Creek, CA 94598, USA 6 (bases 1 to 97083)
DOE Joint Genome Institute and Stanford Human Genome Center.
                                                                                                                                                                                                                         position 96045.
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DOE Joint Genome Institute and Stanford Human Genome Center.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Baldwin, J., Barna, N., Beckerly, R., Beda, F., Boguslavkiy, L., Boukhgalter, B., Brown, A., Burkett, G., Castle, A., Chepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Domino, M., Doyle, M., Fenestor, J., Fitzhugh, W., Forrest, C., Gage, D., Galagan, J., Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L., Karatas, A., Klein, J., Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L., Kornan, S., Lenders, T., Lehczky, J., Levine, R., Lieu, C., Liu, G., Locke, K., Macdonald, P., Marquis, N., McSwan, P., McGurk, A., McKernan, K., Meldrin, J., Meneus, L., Morrow, J., Naylor, J., Norman, C. H., O'Connor, T., O'Donnell, P., Olivar, T. M., Peterson, K., Pierre, N., Pisani, C., Pollara, V., Raymond, C., Riley, R., Rothman, D., Roy, A., Santos, R., Severy, P., Spencer, B., Stange-Thomann, N., Stojanovic, N., Subramanian, A., Talamas, J., Tesfaye, S., Theodore, J., Tirrell, A., and Zody, M., Viel, R., Vo, A., Wu, X., Wyman, D., Ye, W.J., Zimmer, A.
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                                                                                                                                                                                                                                                                                                                                                                                              Submitted (14-DEC-1999) whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Mar 30, 2000 this sequence version replaced gi:6692324.
All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 151620)
Birren,B., Linton,L., Nusbaum,C. and Lander,E.
Homo saplens, clone RP11-13P4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and Zody, M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (bases 1 to 151620)
                  Chemistry: Dye-terminator Big Dye; 100% of re Assembly program: Phrap; version 0.960731
Consensus quality: 14737 bases at least Q40
Consensus quality: 145736 bases at least Q30
Consensus quality: 147339 bases at least Q20
Theory 42.2. 12.2.
                                                                                                                                                                                                    Center project name: L3379
Center clone name: L3_P_4
                                                                                                                                                                                                                                                        Contact: sequence_submissions@genome.wi.mit.edu
                                                                                                                            Sequencing vector: M13; M77815; 100% of reads Chemistry: Dye-terminator Big Dye; 100% of reads
                                                                                                                                                                                                                                                                                                         Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                   Center code: WIBR
                                                                                                                                                                                                                                                                                                                                                     Center: Whitehead Institute/ MIT Center for Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HTGS_PHASE1; HTGS_DRAFT.
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113492 151620: contig of 38129 l
Location/Qualifiers
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1432 2326:
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                         /note="assembly_fragment"
22583. .26540
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15258. .17752
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/db_xref="taxon:9606"
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5 42354: gap of 100 bp
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5 53074: gap of 100 bp
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36501: gap of
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4519: contig of 2093 bp in length
4619: gap of 100 bp
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contig of 2495 k
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Male

bp in length

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NOTE: This is a 'working draft' sequence. It currently consists of 18 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
                                                                                                                                                                                  as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Quality coverage: 4.6 in Q20 bases;
Quality coverage: 5.5 in Q20 bases;
                                                                                                         1331: contig of 1331 bp in length
contig of 895 bp in length of 100 hr
                                                                                                                                                                                                                                                                                                                                                                                                                                                              agarose-fp
sum-of-contigs
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bp in length

length

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REFERENCE
AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                   SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DEFINITION
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                                                      Submitted (31-JUL-2001) DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA On Jul 31, 2001 this sequence version replaced gi:13786333. Draft Sequence Produced by DOE Joint Genome Institute
                                                                                                                                                  Submitted (15-SEP-1999) Production Sequencing Facility, DOE Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 9459 (bases 1 to 188460)

DOE Joint Genome Institute and Stanford Human Genome Center.
                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 180460)
DOE Joint Genome Institute and Stanford Human Genome Center.
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Finishing Completed at Stanford Human Genome Center www-shgc.stanford.edu
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DOE Joint Genome Institute.
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HTG.
                                         www.jgi.doe.gov
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Best Local Similarity
Matches 205; Conserv
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                             303 CAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCG
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WI-11780 G23805
WI-11439 G23701
WI-15863 G24307
SHGC-36487 G30204
SHGC-11618 G14612
363 ACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGC
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                                                                                                                                                                                                                       Patent: WO 0194629-A 1119 13-DEC-2001;
Avalon Pharmaceuticals (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                   Young, P.E., Augustus, M., Carter, K.C., Ebner, R., Horrigan, S., Soppet, D.R. and Weaver, Z. Cancer gene determination and therapeutic screer
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/db_xref="taxon:9606"
/chromosome="16"
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Young,P.E., Augustus,M., Carter,K.C., Ebner,R., Endress,G., Horrigan,S., Soppet,D.R. and Weaver,Z.
Cancer gene determination and therapeutic screening using s
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Sequence 1517 from Patent W00194629.
AX331008
AX331008.1 GI:18121642
                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Avalon Pharmaceuticals (US)
Location/Qualifiers
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Patent: WO 0194629-A 1517 13-DEC-2001;
Avalon Pharmaceuticals (US)
Location/Qualifiers
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Pred. No. 6.5e-40;
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Post-processing: Minimum Match 0%
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4: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2001B.DAT: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

91.1 504 19 89.6 506 19	AAH34835 AAA87730 AAF64012	94.9 655 22 AAS944932 93.3 538 22 AAH98515 93.3 538 22 AAH98548	Match Length DB ID  100.0 534 21 AAA38006	
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34.7 34.7 34.7 34.7	39. 38. 38. 38. 4	50.9 42.0 40.1 40.1 40.1	55.2 55.3	61.3 60.3 60.3 58.0 57.9	87.4 87.2 79.6 77.0 75.9 62.7 62.7
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		21 22 22 22 22	22 22 23 23	22 22 22 24	20 21 21 21 21 21
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## ALIGNMENTS

RESULT 1 AAA38006 CN1244584-A UCK-1; chemotaxis factor; immunocyte; haemopoiesis stimulant; tumour; radiotherapy; chemotherapy; human; ss. UCK-1 nucleotide sequence AAA38006 standard; cDNA; 534 BP. 22-AUG-2000 (first entry) 14-MAY-1999; 16-FEB-2000. Homo sapiens 99CN-0107284

Chemotaxic factor useful for treatment and diagnosis of immunocyte disorders - has immunocyte chemotaxic stimulating factor WPI; 2000-388170/34. . . P-PSDB; AAY98142. Ma D, Han W, Zhang Y;

14-MAY-1999;

99CN-0107284.

(UYBE-) UNIV BEIJING MEDICAL

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Best Local :
                    ulcer; osteoporosis; bone degenerative disorder; periodontal disease; gut protection; lung; liver fibrosis; immune deficiency; infection;
                                                                            ischaemia-reperfusion injury; haematopoiesis; cancer; neuropathy; transgenic animal; Alzheimer's disease; Parkinson's disease; burn; amyotrophic lateral sclerosis; platelet disorder; thrombocytopenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence represents an UCK-1 cDNA sequence encoding a chemotaxis factor polypeptide. The UCK-1 protein exhibits immunocyte chemotaxis activity and a haemopoiesis stimulating effect. The invention relates to UCK proteins, their encoding nucleotide sequences and antibodies and antagonists against the proteins. The nucleotide and protein sequences are useful for the preparation of a composition for the diagnosis and treatment of diseases associated with abnormal immunocyte function and low haemopoiesis function caused by radiotherapy and chemotherapy used to
                                                                                                                                                                                                                           cDNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                  AAS44932 standard;
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                                                                                                                                                              secreted protein; arthritis; Crohn's disease; sepsis; shock;
  combined
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                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                        novel human secretory protein, Seq ID No
                                                                                                                                                                                                                                                                                                                                                                                                  cDNA; 655
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No. 4.6e-107;
  autoimmune
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AGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCG

Matches Query Match Best Local

507;

Conservative

0;

Mismatches

0;

0

Gaps

0;

Similarity

94.9%;

Score 507; Pred. No.

DB T;

Length Indels

.6e-1

Sequence 655 BP; 178

A; 129 C; 181

G; 167

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CC involved in increasing haematopoiesis, stem cell survival, bone growth CC and remodeling. (I), (II) and modulators of (II) are useful for CC prophylaxis or treatment of one or more cancers. (II) is also useful for CC creating transgenic animals useful for studying the in vivo activities of CC (I) induces the proliferation of neural cells and regeneration of nerve CC and brain tissue and is useful for the treatment of central and CC peripheral nervous system diseases and neuropathies, such as Alzheimer's, CC Parkinson's disease, Huntington's disease, and amyotrophic lateral CC sclerosis. In addition, (I) is involved in chemotactic or chemokinetic CC activity, regulation of haematopoiesis and is useful for treating myeloid CC or lymphoid cell disorders, platelet disorders such as thrombocytopenia CC and for regeneration of bone, cartilage, tendon, ligament and/or nerve ctissue growth, and in tissue repair, healing of burns, incisions, CC ulcers, for treating osteoporosis, osteoarthritis, bone degenerative CC disorders, or periodontal disease. Furthermore, (I) is also useful for cperfusion injury in various tissues, various immune deficiencies and cC disorders including severe combined immundeficiency (SCID), bacterial or CC fungal infections, autoimmune disorders e.g. multiple sclerosis, alleratic analytics, disheres mellitus, myasthenia gravis, alleratic
                                               analgesic effects or other pain reducing effects, immunoglobulin like activity and can act as an antigen in a vaccine composition to raise immune response. AAS44920-AAS45295 represent novel human secreted pro
                                                                                                              rheumatoid arthritis, diabetes mellitus, myasthenia gravis, allergic reactions and conditions, such as asthma or other respiratory problems. In addition, (I) affects biorhythms or circadian cycles of rhythms, fertility, metabolism, catabolism, anabolism, storage or elimination of dietary fat, lipid, protein, carbohydrate, vitamins, minerals, provides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to novel isolated human secreted polypeptides (I) and polynucleotides (II). (I) and (II) are useful for treating inflammatory conditions such as arthritis, nephritis, Crohn's disease, ischaemia reperfusion injury, shock, sepsis, immune responses, and is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polypeptides and nucleic acids obtained from cDNA libraries prepared from various human tissues, for diagnosis and treatment cancer, neurological, inflammatory, and autoimmune disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tang
Zhao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
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19-SEP-2000;
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19-MAY-2000;
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fertility; analgesic; pain; antigen; ss.
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)B; AAU28032.
                                   sequences
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Zhang J, Cl
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RESULT 3
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XX ID ID-OCT
XX ID ID ID-OCT
XX ID ID ID ID
XX ID ID ID
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17-JUL-2000; 2000US-0617746
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; horse; tomato; monkey; dog; sea urchin; expressed sequence tag; EST;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnostics; forensic test;
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Isolated polypeptide
                                                                                                                                                  (HYSE-) HYSEQ INC.
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                                                                                            Drmanac RA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGAAGTTTTGTAATTTTATATTACTTTTTAGTTTTGATACTAAGTATTAAACATATTTCTG
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gene mapping; genetic disorder;
nutrition; ss.
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Best Local Similarity
                                                  tomato; monkey; dog; sea urchin; expressed sequence tag; ES diagnostics; forensic test; gene mapping; genetic disorder; biodiversity; gene therapy; nutrition; ss.
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                                                                                                                                                  Human EST-derived coding sequence
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                                                                                              sheep; pig; cow; fruit fly; yeast; hamster; macaque; h: monkev: dog: sea urchin; expressed sequence tag; EST;
                                                                                                                                                                                                                                                                 standard;
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The present invention provides the protein and coding sequences of novel proteins from a variety of organisms, including human, dog, cat, horse, cow, ply, hamster, monkey, macaque, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags (ESTs from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is a cDN of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG
                                                                                                                                                                         CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
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                       AAAAAGAAGTTTTGTAATTTTATATTACTTTTTTAGTTTGATACTAAGTATTAAACATATT
                                                                                    CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
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Pred. No. 2.8e-99;
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17-JUL-2000; 2
03-AUG-2000; 2
15-SEP-2000; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 538 BP; 141 A; 142 C; 115
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Drmanac RA,
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TCTGTATTCTTCCAAAAAAAAA
                                       AAAAAGAAGTTTTGTAATTTTATATTTACTTTTTAGTTTGATACTAAGTATTAAACATATT 494
                                                                                             TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA 434
                                                                                                                                                     TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                                                                                            CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
                                                                                                                                                                                                           CAGTATTCATGCTCATCCTATCTGTGTTTGGCACTGATACCAGAAACCACAACATTGACAG
                                                                                                                                                                                                                                                           TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
                                                                                                                                                                                                                                                                                                                      CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT 194
                        AAAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATT
                                                                               TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
                                                                                                                                     TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                                                                                                                                                   TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
                                                                                                                                                                                                                                                                                                       CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT
                                                                                                                                                                                                                                                                                                                                                               ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                           ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                     GGGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGGGGTGCTCCGCCGCGGTGGCGGTTGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 467; 1275pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; 2000US-0491404.
; 2000US-0617746.
; 2000US-0631451.
; 2000US-0663870.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    research use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ou P, Qian
Zhang J, V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 498.2; DB 2
Pred. No. 2.8e-99;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     xB, Wang
Werhman T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 140 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnostics, raising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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RESULT 5
AAH34835
ID AAH3
AX AAH3
AX AAH3
AX AAH3
AX AAH3
AX AAH3
AX Huma
XXX Huma
XXX Homc
OS Homc
OS Homc
XX Hom
PN WO20
XX 29-5
PN 05-A
XX 29-5
PR 03-N
XX WPI;
PA (HUMA
XX HUMA
XX HUMA
XX HUMA
XX CLai
XX WPI;
DR P-PS
XX P-PS
XX CLai
XX NAH:
CC CAAH:
CC
                                                           QΥ
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                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P expression. For example, N and P may be used to treat disorders associated with inappropriate P expression for example, N and P may be used to treat disorders or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention. N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding 4277 human useful for preventing, diagnosing
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 558 BP; 150 A; 119 C; 145 G; 141 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 3428; 9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-235357/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ruben SM, Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200122920-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human colon cancer antigen encoding cDNA SEQ ID NO:1917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH34835 standard; cDNA; 558 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                            499;
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                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAG75430.
                                                                                                                                                               TGAAGCCGAGCTGGGCGAGAAGTAGGGGGGGGGGGGGCGCTCCGCCGCGGTGGCGGTTGCTA
TCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGC
                                                                                                                              TGAAGYSGGAGCTGGCGAGAAGTAGGGGAGGGCGGTGCCTCCGCCGCGGTGGCGGTTGCTA
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                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000WO-US26524.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         990S-0157137.
990S-0163280.
                                                                                                                                                                                                                                                                                   93.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4277 human colon cancer-associated polypeptides, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Birse CE,
                                                                                                                                                                                                                                                            2,
                                                                                                                                                                                                                                                                                      Score 497.2; DB 2
Pred. No. 4.7e-99;
                                                                                                                                                                                                                                                            Mismatches
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RESULT 6
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AAAABTTA
AAAATTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cystic fibrosis; hypothyroidism; immunological disorder; amyloidosis; brain disorder; skeletal muscle disorder; eye disorder; obesity; mitochondriocytopathy; diabetes; atherosclerosis; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted protein encoding cDNA SEQ ID #29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA87730;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA87730 standard; cDNA; 515
  This sequence represents human cDNA encoding a secreted protein. Invention relates to sequences AAA87725-A87774 which encode human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200037491-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-NOV-2000
                                                                                   Claim 1; Page 169-170; 306pp; English
                                                                                                                                                                Polynucleotides and polypeptides encoding proteins with signeptides, useful in diagnostic, forensic, gene therapy and
                                                                                                                                                                                                                                                                                                                                    Bougueleret L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-DEC-1998;
25-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; secreted protein; forensic procedure; gene therapy;
                                                                                                                                                                                                                                                      P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                          (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neurodegenerative disorder; graft rejection; dementia; hyperlipidaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chromosome mapping; cancer; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    491
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DB; AAB25768.
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                                                                                                                                         procedures
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTGTATTCTTCCAAAAAAAAAAAAAAA 522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAAAGAAGTTTTGTAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTT 495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGTATTCATGCTCATCGTATCTGTGTTTGGCACTGATACCAGAAACCACAACATTGACAGT
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99US-0141032.
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                                                                                                                                                                                                                                                                                                                                          Duclert
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                                                                                                                                                                                                signal
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RESULT 7
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AC AAF6
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KW Secr

05-APR-2001

(first entry)

AAF64012 standard; cDNA; 515

ВP

Secreted protein; prevention; treatment; diagnosis; disease;

cDNA encoding human secreted protein #13.

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The cDNAs are useful for expressing secreted proteins or fragments to obtain antibodies capable of specifically binding to the secreted proteins. The cDNAs may also be useful in diagnostic, forensic, gene therapy and chromosome mapping procedures and may be used to design expression vectors and secretion vectors. The proteins of the invention may be used to treat diseases including cancer, autoimmune diseases, cardiovascular diseases, exist fibrosis, hypothyroidism, immunological disorders, amyloidosis, brain disorders, skeletal muscle disorders, eye disorders, obesity, mitochondriocytopathies, diabetes, atherosclerosis, neurodegenerative disorders, graft rejection, Alzheimer's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 secreted proteins AAB25763-B25812. The proteins include signal peptides. Included in the invention are a host cell containing one of the cDNA sequences, and a purified antibody capable of binding to one of the secreted proteins. Also contained in the invention are methods for storing the sequence data on a computer system, and a method for identifying features of the cDNA sequences using a computer programme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 515 BP; 143 A; 106 C; 135 G; 130 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       dementia, hyperlipidaemia, septic shock and impotence.
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                                                                                                                                                                                                                                                                                                                                                        CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAGAGTTGAGGGAAAGTGCTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGGAAGCCGAGCTGGGCGAGAAGTAGGGGGAGGGCGGTGCTCCG-CGCGGTGGCNGTTGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTGAAGCCGAGCTGGGCGAGAAGTAGGGGAAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT
                                                                                         AAAAGAAGTTTTGTAATTTTATATTTACTTTTTAGTTTGATACTAAGTATTAAACATATT
                                                                                                                                                             TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
                                                                                                                                                                                                                   TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                                                                                                                                                 TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                                                                                                                                                                                                          CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
                                   TCTGTATTCTTCCAAAAAAAAAAAAAA 522
                                                                                                                                            TTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAA
 TCTGTATTCTTCCAAAAAAAAAAAAAA 514
                                                                     AAAAAGAAGTTTTGTAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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Pred. No. 3.1e-98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                             Sequence 515
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 225; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene
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21-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Secreted proteins and the nucleic acids encoding ne therapy and for detecting similar sequences in
                                                                       367
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                                                                                                                                                                                                                       CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT 194
                                TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA 374
       TCTGTATTCTTCCAAAAAAAAAAAAAAAA
                                                                                                         TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCCTTA
                                                                                                                                               CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
                                                                                                                                                          CAGTATTCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAG
                                                                                                                                                                                    TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
                                                                                                                                                                                            TCAGTGTGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAA
                                                                                                                                                                                                                                                            GTGAAGCCGAGCTGGGCGAGAAGTAGGGGGAGGGCGGTGCTCCGCCGCGCGGTGGCGGTTGCT
                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                           BP; 143 A; 106 C; 135 G; 130 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  990S-0141032
990S-0469099
                                                                                                                                                                                                                                                                                                                                                 92.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bougueleret L,
                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                 Score 493.4; DB Pred. No. 3.1e-98
                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                               ..1e-98;
                                                                                                                                                                                                                                                                                                                                                         22; Length
                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                           them, useful samples -
                                                                                                                                                                                                                                                                                                                                                         515;
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                                                                                                                                                                                                                                                                                                                                                                                                                          CDNAS
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RESULT 8

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Human; secreted protein; fusion protein; gene therapy; protein therapy; dlagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; inflammation; ischaemic shock; Alzheimer; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer; disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
11-SEP-1998
                                                                                      W09839448-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted protein gene 88 clone HAUAV32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV59598 standard; DNA; 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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11 - APR - 1997;
23 - MAY - 1997; 07-MAR-1997; 07-MAR-1997; 07-MAR-1997; 07-MAR-1997; 07-MAR-1997; 02-OCT-1997; 07-MAR-1997; 07-MAR-1997; 06-MAR-1998; 97US-0040626 97US-0043311 97US-0043313 97US-0043569 97US-0043576 97US-0043576 97US-0043570 97US-0043671 97US-0043671 97US-0043671 97US-0047501 97US-0047501 97US-0047501 97US-0047581 97US-0047581 97US-0047581 97US-0047582 97US-0047583 97US-0047583 97US-0047583 97US-0047583 97US-0047583 97US-0047583 97US-0047583 97US-0047587 97US-0047593 97US-0047593 97US-0047593 97US-0047593 97US-0047598 97US-0047598 97US-0047598 97US-0047598 97US-0047598 97US-0047598 97US-0047599 97US-0047598 97US-0047599 97US-0047599 97US-0047599 97US-0047599 98WO-US04493. 97US-0040161

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23 -MAY -1997

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27 -MU
This sequence represents a nucleic acid molecule designated Gene 88 from the human cDNA clone HAUAV32 (deposited as clone ATCC 97901 and ATCC 209047) which encodes a secreted human protein. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin Fc portion (e.g. AAV59502) for increasing the stability of
                                                                                                                                                     New isolated human genes and the secreted polypeptide(s) they encode - useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
                                                                                                               Claim 1; Page 320; 721pp; English.
                                                                                                                                                                                                                                                                                                                                               Bednarik DP,
                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN
                                                                                                                                                                                                                                                                         P, Ferrie n., Li Y, MG
H, Lafleur DW, Li Y, MG
an SM, Shi Y, Soppet DR,
                                                                                                                                                                                                                               1998-506364/43.
DB; AAW74818.
                                                                                                                                                                                                                                                                                       DP, Brewer LA, Carter KC, Duan R, Ebner R, El
Ferrie AM, Fischer CL, Florence KA, Greene JM,
Lafleur DW, Li Y, Moore PA, Ni J, Olsen HS, J;
Shi Y, Soppet DR, Young PE, Yu GL, Zeng Z;
                                                                                                                                                                                                                                                                                                                                                                                      GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-0047618

97US-0047633

97US-0048964

97US-0048974

97US-0051926

97US-0055724

97US-0056630

97US-0056631

97US-0056631

97US-0056635

97US-0056637

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97US-0056845

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97US-0056879

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97US-0056889

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97US-00568881
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97US-0047613.
97US-0047614.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the fused protein as compared to the human protein only. The invention relates to 186 novel genes and their fragments (nucleic acid sequences: AAV59511-V59812; amino acid sequences AAW74731-W75026) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypeptides in a sample or by determining the presence of mutations in the new polynucleotides. Specific uses are described for each of the 186 polynucleotides, based on which tissues they are most highly expressed in polynucleotides, based on which tissues they are most highly expressed in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (see AAV59511 for described uses).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 CCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                    CTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGCTG
                                                                                                                                                                                                                                                                                                    TCATGCTCATCGTATCTGTTGTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTG
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                                                                                                                                                             GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC
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                                 TTCTTCCAAAAAAAAAAAAAAAAA 525
                                                                AAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTA
                                                                                      AAGTTTTGTAATTTTATATTACTTTTTAGTTTTGATACTAAGTATTAAACATATTTCTGTA
                                                                                                                                          GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC
                                                                                                                                                                                                                                                                                   TCATGCTCATCGTATCTGTGTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTG
TTCTTCCAAAAAAAAAAAAAAAAA 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               143 A; 106 C; 128 G; 127 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 486.6; DB 19; Length Pred. No. 9.4e-97;
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AAV59746 standard;

(first entry)

Human secreted protein gene 88 clone HAUAV32.

AAV59745
AAV59745
AAV59745
AC AAV5
XX AAV5
XX AAV5
XX AAV5
XX Huma
XX Huma
KW diag
KW deve
KW diag
KW deve
KW inmu
KW inmu
KW inmu
KW inmu
KW ingu
KW cogu Human; secreted protein; fusion protein; gené therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; ALDS; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm

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11-APR-1997
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23-MAY-1997
23-MAY
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970S-0047596.
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970S-0047600.
970S-0047601.
970S-0047611.
970S-0047613.
           97US-0052874.
97US-0055724.
97US-0056630.
97US-0056631.
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97US-0047617.

97US-0047618.

97US-0047632.

97US-004963.

97US-0048964.

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23-AUG-1997
24-AUG-1997
25-SEP-1997
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25-SEP-1997
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22-AUG-1997;
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22-AUG-1997;
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22-AUG-1997;
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97US-0056882.
97US-0056884.
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97US-0056637.
97US-0056662.
97US-0056664.
97US-0056845.
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Bednarik DP, Brewer LA, Carter KC, Duan R, Ebner R, Endress Feng P, Ferrie AM, Fischer CL, Florence KA, Greene JM, Hu J Kyaw H, Lafleur DW, Li Y, Moore PA, Ni J, Olsen HS, Rosen Ruben SM, Shi Y, Soppet DR, Young PE, Yu GL, Zeng Z; Endress GA; JM, Hu JS; , Rosen CA;

P-PSDB; AAW74961. WPI; 1998-506364/43.

New isolated human genes and the secreted polypeptide(s) they encode - useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders

Claim 1; Page 472; 721pp; English.

This sequence represents a nucleic acid molecule designated Gene 88 from C the human cDNA clone HADAV32 (deposited as clone ATCC 97897 and ATCC 209043) which encodes a secreted human protein. The gene can be used to generate fusion proteins by linking to the gene to a human C immunoglobulin Fc portion (e.g. AAV59502) for increasing the stability of the fused protein as compared to the human protein only.

C The invention relates to 186 novel genes and their fragments (nucleic acid sequences: AAV59511-V59812; amino acid sequences AAW74731-W75026) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new conjuncteotides. Specific uses are described for each of the 186 polynucleotides, based on which tissues they are most highly expressed in (see AAV59511 for described uses).

Sequence 506 BP; 148 A; 103 C; 121 G; 132 T; 2 other;

Query Match Best Local Similarity Matches 484; Conserv Conservative 89.6%; 97.4%; Score 478.6; DB 19; Length 506; Pred. No. 5.2e-95; Mismatches 11; Indels 0; Gaps 0

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RESULT 10
AAX9726
ID AAX97
XX AAX97
XX 23-SE
XX Secre
KW Glagn
XX W0992
XX W0992
XX W0992
XX W0992
XX W0992
XX 04-SE
PR 13-NC
XX 04-SE
PR 13-NC
XX 04-SE
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17-DEC-1997;
09-FEB-1998;
13-APR-1998;
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cDNAs encoding
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97US-0066677.
97US-0069957.
98US-0074121.
98US-0081563.
98US-0096116.
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XX 12-JU
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DX Human
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KW Cytok
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97.8%;
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Pred. No. 1.9e-92;
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cytokine production; cell proliferation; cell differentiation; immune deficiency; infectious disease; autoimmune disorder; asthma; multiple sclerosis; systemic lupus erythematosus; rheumatoid arthritis; allergic reaction; osteoporosis; osteoarthritis; periodontal disease; nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; liver fibrosis; lung fibrosis; reperfusion injury; systemic cytokine damage; tissue differentiation; contraceptive; stroke; coagulation disorder; myoccardial infarction; inflammatory condition;

Human

protein clone HP10357 full length coding sequence.

hydrophobic domain; nutritional source; haematopolesis;

protein;

12-JUN-2000

(first entry)

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Conditions of the treatment of suppressing activity. It can be used in the conditions. The protein is also used for treatment of various immune deficiencies and disorders, and to treat infectious diseases caused by viral, bacterial, fungal or other conditions. The protein is also used for treating autoimmune disorders conditions and trible sclerosis, systemic lupus erythematosus, and rheumatoid conditions such as asthma, and in immune suppression after organ consequently in the treatment of allergic reactions and consequently in the treatment of myeloid or lymphoid cell cand consequently in the treatment of myeloid or lymphoid cell conditions in the protein is useful in regulation of haematopoiesis deficiencies. It is also used in compositions for tissue growth or regeneration. The protein is also used in the treatment of steoporosis cor osteoarthritis and in the treatment of periodontal disease and other tooth repair processes. The protein is used in the treatment of nervous consistent disorders such as Alzheimer's disease, Parkinson's disease, and conting or liver fibrosis, reperfusion injury in various consistent of lung or liver fibrosis, reperfusion injury in various calso used for promotting or inhibiting tissue differentiation. They are calso used as contraceptives since they exhibit activit or inhibit related creatment and prevention of creatme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-JUL-1998;
07-AUG-1998;
25-AUG-1998;
09-SEP-1998;
29-SEP-1998;
                                                           conditions resulting from coagulation activities e.g. myocardial infarction or stroke. They also acts as receptors, receptor ligands or inhibitors or agonists of receptor/ligand interactions. They are used to treat inflammatory conditions such as septic shock, sepsis, ischaemia reperfusion injury, arthritis, and nephritis. They can be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nutritional sources or supplements. The protein exhibits cytokine, cell proliferation, cell differentiation activities and induces production of other cytokines in certain cell populations. The protein also exhibits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence encodes a human protein of the invention, which has hydrophobic domains. The DNA sequences can be used as a probe or as a genetic marker. The protein can also be used as a marker, and to identify potential genetic disorders. The DNA and protein can also be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human proteins having hydrophobic domains useful for treating osteoporosis, Alzheimer's disease, Parkinson's disease, asthma, multiple sclerosis, rheumatoid arthritis, cancer, anaemia, and stro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 228; 351pp; English.
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98JP-0224105.
98JP-0238116.
98JP-0254736.
98JP-0275505.
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                                                                                                             Merberg D,
Clark HF,
                                                                                       WPI; 2002-179322/23.
                                                                                                                                     Jacobs K,
                                                                                                                                                                                                                                                                                                    Human secreted protein; hyperproliferative disorder; autoimmune disorder; immune deficiency disorder; blood disorder; inflammatory disorder; infectious disorder; allergic condition; neurodegenerative disorder; liver fibrosis; coagulation disorder; gene therapy; antimicrobial;
                                                                                                                                                                                 06-APR-2000; 2000US-195605F
                                                                                                                                                                                                                               18-OCT-2001.
                                                                                                                                                                                                       29-MAR-2001; 2001WO-US10232
                                                                                                                                                                                                                                                     WO200177289-A2
                                                                                                                                                                                                                                                                                                                                                                 cDNA sequence #397 encoding novel human secreted protein.
                                                                                                                                                                                                                                                                                                                                                                                           08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK36006 standard; cDNA; 439
                                                                                                                                                          (GEMY ) GENETICS INST
                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                 cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTACTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTT
                                                                                                           McCoy JM,
Treacy M,
Fechtel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTACTTTTTAGTTTTGATACTAAGTATTAAACATATTTCTGTATTCTT 505
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATC
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                                                                                                                                                                                                                                                                                               hepatotropic;
                                                                                                                                   Lavallie ER,
                                                                                                                                                           INC.
                                                                                                                      Agostino MJ,
                                                                                                              Howes
                                                                                                             SH, Resnick RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                              immunosuppressive;
                                                                                                          Collins-Racie LA, Evar
, Bowman MR, Spaulding
Resnick RJ, Gulukota K,
                                                                                                                                Evans C;
                                                                                                                       <
                                                                                                           Graham
                                                                                                                      Wong
                                                                                                                                                                                                                                                                                             gene;
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Matches Query Match

Госат

Similarity

87.2%; 99.8%; A; 101

Score 465.4; DB 2 Pred. No. 3.8e-92; 0; Mismatches 1

Indels

0;

Gaps

0

Claim 1; Page 296-297; 393pp; English

Six hundred and twenty three polynucleotides derived human tissue sources which encode secreted proteins.

es derived from a variety of proteins, useful for

disorders

treating immune deficiencies and disorders such as autoimmune

Sequence 467 BP; 118

C; 121 G; 127

T; 0 other DB 21; Length 1:

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88888888888888888888888888888888888
RESULT 13
AAX97873
ID AAX97
XX AAX97
AC AAX97
XX 23-SE
DT 23-SE
XX Humar
XX Secre
KW Glagr
XX dlagr
XX Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC The present invention relates to the isolation of novel cDNA sequences cc which encode human secreted proteins. The cDNA sequences have been cc derived from a variety of human tissues. The invention also provides can method for producing proteins from these polynucleotide sequences. CC The proteins are useful for identifying compounds that modulate their cativity and production. The sequences of the invention are useful for treating diseases such as hyperproliferative disorders cc (e.g. cancer), immune deficiency disorders (e.g. severe combined cc immunodeficiency (SCID)), autoimmune disorders (e.g. multiple cc sclerosis), blood disorders (e.g. thrombocytopaenia), inflammatory cc disorders (e.g. arthritis), infectious disorders (e.g. hepatitis), cc allergic conditions (e.g. asthma), neurodegenerative disorders (e.g. allergic conditions (e.g. asthma), neurodegenerative disorders (e.g. C Alzheimer's disease), liver fibrosis, coagulation disorders (e.g. cc invention are also useful in gene therapy. ABK35610-ABK36232 represent cc servered morefules.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 439 BP; 117 A;
                                                                    Secreted protein; human; cytostatic; diagnostic; gene therapy; chromosome
                                                                                                                                                                                               AAX97873;
                                                                                                                                                                                                                                 AAX97873 standard; cDNA; 413
                                                                                                                       Human secreted protein encoding cDNA
                                                                                                                                                           23-SEP-1999
                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                      421
                                                                                                                                                                                                                                                                                                                                                         491
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                      TATTTCTGTATTCTTCC 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTGCTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTGCTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAAACATCGCCCCTTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TATTICTGTATTCTTCC 507
                                                                                                                                                                                                                                                                                                                                                                                                                           AAAAAAAAGAAGTTTTGTAATTTTATATTACTTTTAGTTT-GATACTAAGTATTAAACA 490
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAAGCCTGTGCATG 431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                      gene therapy; chromosome mapping; secretion vector; ss.
                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          79.6%;
99.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              96 C; 104 G; 122 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 425; DB 24;
Pred. No. 2.3e-83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                      thrombotic; osteopathic; forensic;
                                                                                                                           #61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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W09925825-A2

RESULT 14
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06-OCT-2000 AAC00147; AAC00147

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Matches
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13-NOV-1997;
17-DEC-1997;
09-FEB-1998;
13-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                              AAX97813-X97906 represent extended cDNA's which encode novel human secreted proteins (see AAY36129-Y36222) and which have cytostatic, thrombotic and osteopathic activity. The extended cDNAs can be used texpress secreted proteins or parts of them or to obtain antibodies capable of binding to the secreted proteins. They may also be used in diagnostic, forensic, gene therapy and chromosome mapping procedures.
                                                                                                                                                                                                                                                                                                                                                                                                              diagnostic, forensic, gene therapy and chromosome mapping processors also include design of expression vectors and secretion vectors
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                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 261; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Extended cDNAs encoding secreted proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; AAY36189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEST )
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                                                                                                                                                                                                                                                                                                                                              412;
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                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                             GTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGT
                                                                                                                                                        CTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCA 346
                                                                                                                                                                                                         CTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTTGGCA 286
                                                                                                                                                                                                                                                CCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAAGGCCACGTGAAGATGCTGCGG
                                                                                                                                                                                                                                                                      CCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCCACGTGAAGATGCTGCGG
                                                                                                                                                                                                                                                                                                                                                                                               413 BP; 121 A; 82 C; 95 G; 115 T; 0 other;
CCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAGAAGTTTTGTAATTTTATATTTACTTTT
                                                          CCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAGTTTTTGTAATTTTATATTACTTTT
                                                                                              CTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGCA
                                                                                                                                                                                               CTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTGGCA
                                                                                                                                                                                                                                                                                                                                              Conservative
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98US-0074121
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97US-0066677
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99.8%;
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Best Local Similarity
Matches 406; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5' ESTs were prepared from the sequence of polya+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 415 BP; 101 A; 97 C; 126 G; 91 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID 145; 71pp + CD-ROM; English.
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                                                    ATCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTG 134
                                                                                                              CTGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCT 194
                                                                                                                                                                                                                                                                                                          GTGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCT 74
               TIGGIGGAGGGGIGITIGCACITGIGACAGCAGIAIGCIGICTIGCCGACGGGGCCCTIA 374
                                                                                                                                                                                                                                                                                         GGGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCCTCCGCCGCGGTGGCGGTTGCT 68
TTGGTGGAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTA
                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                Score 405.4; DB 21; Length 415; Pred. No. 4.2e-79;
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23-APR-1998
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13-MAR-1998;
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17-MAR-1998;
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11-MAR-1998;
11-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PR0772 nucleotide sequence
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98US-0079663

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98US-0079728

98US-0079729

98US-0079729

98US-0080105

98US-0080107

98US-0080107

98US-0080327

98US-0080334

98US-0080334

98US-0081071

98US-00811071

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98US-008120

98US-0081256

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98US-0082704
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98US-0079294.
98US-0079656.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transmembrane protein; ss.
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07-MAY 1998
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13-MAY 1998
11-MAY 1998
                             The present invention describes secreted and transmembrane polypeptides and their polynuclectides. The nucleotide sequences are useful as sources of probes, primers, for chromosome mapping, and for generation of antisense sequences. They can also be used to create transgenic animals. The proteins can be used to treat a variety of diseases and disorders, depending on their function. Diseases that may be treated include blood coagulation disorders, cancers and cellular adhesion disorders. They may also be used to raise antibodies. AAZ3891 to AAZ34338, and AAY41685 to AAY41774 represent polynucleotide and polypeptide sequence given in the exemplification of the present
                                                                                                                                                                           Claim 2; Fig 69; 530pp; English.
                                                                                                                                                                                                       adhesion
                                                                                                                                                                                                              New secreted and transmembrane polypeptides and their polynucleotides, useful for treating blood coagulation disorders, cancers and cellular
                                                                                                                                                                                                                                                                                                Wood
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P-PSDB; AAY41713.
                                                                                                                                                                                                                                                                                                                       (GETH )
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                                                                                                                                                                                                       disorders
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98US-0085700.

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98US-0086073.

98US-0086414.

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                                                                                AGCAGTATGCTGTCTTGCCGACGGGGCCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAG
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Search completed: June 25, Job time : 219 secs

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Query Match Best Local Similarity

62.7%; 76.0%; A; 137

Score 335; DB 20; Pred. No. 9.8e-64;

Length 663;

Sequence 663 BP; 180

C; 153

G; 193 T; 0 other;

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Title:
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1,

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3: /cgn2_6/ptodata/1,

4: /cgn2_6/ptodata/1,

5: /cgn2_6/ptodata/1,

6: /cgn2_6/ptodata/1,
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534
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Listing first 45 summaries
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US-09-149-476-252
US-09-345-96-10
US-08-545-196B-12
US-08-865-297-5
US-08-865-297-5
US-08-865-297-6
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US-08-95-749-2
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Sequence 98, Appl
Sequence 95, Appl
Sequence 10, Appl
Sequence 11, Appl
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29 41.8 7.8 742 1 US-07-847-010-12 Sequence 12, Appli 31 41.2 7.7 2671 6 5168051-9 Patent No. 5168051 32 41.2 7.7 3459 2 US-08-980-060-3 Sequence 3, Appli 32 41.2 7.7 3459 4 US-09-800-60-3 Sequence 3, Appli 34 41 7.7 3459 4 US-09-800-60-3 Sequence 3, Appli 35 40.8 7.6 1046 1 US-08-481-981-8 Sequence 4, Appli 36 40.8 7.6 1046 1 US-08-481-458-2 Sequence 4, Appli 37 40.6 7.6 2444 3 US-08-481-458-2 Sequence 4, Appli 38 40.6 7.6 2444 3 US-08-451-458-2 Sequence 18, Appli 39 40.6 7.6 2625 4 US-09-245-041-18 Sequence 18, Appli 40.4 7.6 1253 2 US-08-984-919A-10 Sequence 19, Appli 40.4 7.6 1466 4 US-08-984-919A-10 Sequence 10, Appli 40.4 7.6 1466 4 US-08-984-919A-12 Sequence 10, Appli 40.4 7.6 1472 4 US-08-984-919A-12 Sequence 10, Appli 40.4 7.6 1472 4 US-08-984-919A-12 Sequence 10, Appli 41 40.4 7.6 1472 4 US-08-984-919A-12 Sequence 10, Appli 41 41 41 41 41 41 41 41 41 41 41 41 41	9476  9476  ecreted proteins  09/149,476  /US98/04493  040,162  040,333  038,621  040,626  040,336  040,626  040,336  047,600  047,597	tion US/09149476  al.  al.  186 Human Secre 102P1  1098-09-08  11998-03-06  11998-03-06  11998-03-06  11997-03-07  100MBER: 60/040,  11997-05-23  100MBER: 60/047,  11997-05-23	6-98  6-98  6-98  6-98  6-98  6-98  6-98  FORMATION:  RENCE: PCO REPLICATION  TLING DATE REPLICATION  TILING DATE  TILING DATE	SULT 1 -09-149 -09-149 GENERAL APPLICE FILE R FILE R FILE R FILE R FARLIE EARLIE EARLI	A D トラストラストラストライト・アイト・アイト・アイト・アイト・アイト・アイト・アイト・アイト・アイト・ア
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FILING DATE: APPLICATION NUMBER: 60/043,672 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,313 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60 FILING DATE: 1997-04-1 APPLICATION NUMBER: 60/ FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/ FILING DATE: 1997-04-11 FILING DATE: 1997-05-23
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60/047,633

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1997-03-07

60/047,600

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GENERAL INFORMATION:
APPLICANT: ROSED et al.
TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P1
CURRENT APPLICATION NUMBER: US/09/149,476
CURRENT FILING DATE: 1998-09-08
EARLIER APPLICATION NUMBER: PCT/US98/04493
EARLIER FILING DATE: 1998-03-06
EARLIER FILING DATE: 1997-03-07
EARLIER FILING DATE: 1997-03-07
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ER APPLICATION NUMBER: 60/049,610
ER FILLING DATE: 1997-06-13
ER APPLICATION NUMBER: 60/061,060
ER FILING DATE: 1997-10-02
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Pred. No. 2.7e-115;
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ER APPLICATION NUMBER: 60/056,631
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,845
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,892
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ER APPLICATION NUMBER: 60/057,761
ER FILING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/047,595
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ER FILING DATE: 1997-06-06
ER APPLICATION NUMBER: 60/057,650
ER FILING DATE: 1997-09-05
ER APPLICATION NUMBER: 60/056,884
ER FILING DATE: 1997-08-26
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                                                                                                                                                                                                                                                      GATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATC
                                                                                                                                                                                                                                                                                                        TAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCACGTGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1997-09-05
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498
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 478.6; DB 4;
Pred. No. 3e-113;
2; Mismatches 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
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Gaps

96 0

181

216

276

121 156 61

336 241

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; NAME/KEY: misc_feature
; LCCATION: (1)...(392)
; OTHER INFORMATION: n = A,T,C or G
US-09-385-982-95
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: ENDEGE, WILSON O., ET AL.
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS: II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 95, Appl Patent No. 626233
                                                             Patent NO. DOWNSTION:
GENERAL INFORMATION:
APPLICANT: MELKI, JUDITH
APPLICANT: MIUNICH, ARNOLD
TITLE OF INVENTION: SURVIVAL MOTOR NEURON (SMN) GENE:
TITLE OF INVENTION: FOR SPINAL MUSCULAR ATROPHY
TITLE OF INVENTION: 65
                                                                                                                                                                                                             Sequence 10, Application US/08545196B Patent No. 6080577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 297; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EARLIER APPLICATION NUMBER: 09/328,111
EARLIER FILING DATE: 1999-06-08
EARLIER APPLICATION NUMBER: 60/117,393
EARLIER FILING DATE: 1999-01-27
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER FILING DATE: 1998-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/385,982 CURRENT FILING DATE: 1999-08-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: BIRCH, STEWART,
STREET: PO BOX 747
STREET: PO TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         286 ACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGC 345
                                                                                                                                                                                                                                                                                                                                                                       524 AAAAAAAAAA 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                     103 ACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGACAGC 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        226 GCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTTGGC
                                                                                                                                                                                                                                                                                                                                  343 TNAANTTANAA 353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAAAAGTTTTGTAATTTTATATTACTT 282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCCTTACCAGAAAAAGCCTGTGCATG-AAAAAAAAGAAGTTTTGTAATTTTATATTACTT 464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGTATGCTGTCTTGCCGACGGGGCCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGG
                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09385982
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                                                           KOLASCH AND BIRCH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 392;
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US-08-545-196B-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: MELKI, JUDITH
APPLICANT: MUNICH, AROLD
APPLICANT: MUNICH, AROLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY AGENT INFORMATION:

NAME: FARACI, C. J.

REGISTRATION NUMBER: 32,350
REFERENCE/DOCKET NUMBER: 2121
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 205-8000
TELEFAX: (703) 205-8050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (703) 205-8050 INFORMATION FOR SEQ ID NO: [
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                                                                    FILING DATE: 19-OCT-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: FARACI, C. J.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
ADDRESSEE: BIRCH, STEWART, KOLASCH AND BIRCH, LLP
STREET: PO BOX 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                               REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: SURVIVAL MOTOR NEURON (SMN) GENE: A GENE TITLE OF INVENTION: FOR SPINAL MUSCULAR ATROPHY
                                                                                                                                                                                              COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Balcor-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: double TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/0 FILING DATE: 19-OCT-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 9.4%;
Local Similarity 71.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                         CITY:
                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                         STREET:
               TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1487
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            443 GITTTGTAATTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATT 502
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(703) 205-80
03) 205-8050
                                                                                                                                           19-0CT-1995
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                                                                      32,350
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Pred. No. 0.00078;
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                                                        2121-110P
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                                                                                                                                                                                                        Version #1.30
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    Matches
                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                     TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 770 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                   NAME: Silveri, Jean M.
REGISTRATION NUMBER: 39,030
REFERENCE/DOCKET NUMBER: DEN-024
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/OFFILING DATE: 29-MAY-1997 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                               FEATURE:
                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: The Siva Genes, No. 6010853el Genes Involved in TITLE OF INVENTION: CD27-Mediated Apoptosis
                                                                              NAME/KEY:
                Local Similarity
                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Boston
                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1427 GTACTGTTTTTTTCTATCTTCTATATGTTTAAAAGTATATAAAAAAATATTTAATTTTT 1486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           503 CTTCCAAAAAAAAAAAAAAAAAAAAAAAAA 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      66;
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                                                                                                                                                                         nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1582 base pairs
                                                                                                                                                                                                                                     : (617)227-7400
(617)227-5941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
    Conservative
                                                                             CDS
88..612
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                             linear
                                                                                                                              CDNA
                                                                                                                                                          single
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71.78;
                8.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                            us/08/865,297
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Release #1.0, Version #1.25
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Score 46.8; DB 3;
Pred. No. 0.0048;
0; Mismatches 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
 47;
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                             Length 770;
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 Indels
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RESULT 8
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;Patent No. 5194596
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US-08-890-865A-2
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                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212)391-052
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 10-JUL-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 28,678
                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Cooper & Dunham LLP STREET: 1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                  3662 AATGACTTGTACAATTATCTTTTCAAAGGTACTTGGATAATAATGAAATAAAAACGTTTT 3721
                                                                                                    438 AAGAAGTTTTGTAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  533 AA 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  628 CAGCACAAGATGTTCACACTAAAGAGAGAGAGAGAGCTTTTTTATATGTTATGTTTTATA 687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     413 CAGAAAAAGCCTGTGCATGAAAAAAAAGAAGTTTTGTAATTTTATATTTACTTTTTAGTTT 472
                                                                                                                                                                                                                      65;
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                                                                                  TGAACCTTCCAAAAAAAAAAAAAAAAAAAAAAAAAA 3758
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                                                                                                                                                                                                                                                                                                                                                                  3761 base pairs
                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                                                                                                                                                                    8.6%;
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                                                                                                                                                                                                                                  Score 45.8; DB Pred. No. 0.016;
                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                 DB 4; Length 3761;
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                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                497
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;C.;MITCHELL, RICHARD

APPLICANT:

TISCHER,

EDMUND G.; ABRAHAM, JUDITH A.; FIDDES, JOHN

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5219739-16
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                                                                                                                                                                                                          RESULT 10
US-09-276-400-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5219739-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 27-JUL-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JOHN C.; MITCHELL, RICHARD L.
TITLE OF INVENTION: DNA SEQUENCES ENCODING BYEGF120 AND
HYEGF 121 AND METHODS FOR THE PRODUCTION OF BOVINE AND HUMAN
VAASCULAR ENDOTHELIAL CELL GROWTH FACTORS, BYEGF120 AND HYEGF121
                                                                                                                                              Sequence 1, Application US/09276400 Patent No. 6140056 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches 69; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
              CURRENT APPLICATION NUMBER: US/09/276,400 CURRENT FILING DATE: 1999-03-25 NUMBER OF SEQ ID NOS: 10
                                                                        TITLE OF INVENTION: NOVEL MSP-18 PROTEIN AND NUCLEIC ACID MOLECULES AND TITLE OF INVENTION: USES THEREFOR FILE REFERENCE: MNI-073
                                                                                                                                  APPLICANT: Khodadoust, Mehran
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 387,545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: TISCHER, EDMUND G.; ABRAHAM, JUDITH A.; FIDDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 32
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: PRODUCTION OF VASCULAR ENDOTHELIAL CELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 27-JUL-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 450,883 FILING DATE: 14-DEC-1989 APPLICATION NUMBER: 387,545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/07/559,041 FILING DATE: 27-JUL-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            426 TGCATGAAAAAAAGAAGTTTTGTAATTTTATATTACTTTTAGTTTTGATACTAAGTATT 485
                                                                                                                                                                                                                                                                                      426 TGCATGAAAAAAAAGAAGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATT 485
                                                                                                                                                                                                                                                                                                                          486 АЛАСАТАТТТСТБТАТТСТТССАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛАЛ 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  69;
                                                                                                                                                                                                                                                                                                                                                                                                                                          8.4%; ilarity 63.3%; Conservative
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63.3%;
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    Mismatches

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Pred. No. 0.015;
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; LOCATION: (143)..(2401)
US-09-276-400-1
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US-09-448-076-1
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LENGTH: 2920
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LENGTH: 2920
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Best Local Similarity 68.1%;
Matches 62; Conservative
                                                                                                                             GENERAL INFORMATION:
APPLICANT: Khodadoust, Mehran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Applic Patent No. 6300092
                                                                                                                                                                 Sequence 1, Application US/09702572 Patent No. 6391602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/448,076
CURRENT FILING DATE: 1999-11-23
EARLIER APPLICATION NUMBER: 60/117,580
EARLIER FILING DATE: 1999-01-27
EARLIER APPLICATION NUMBER: 09/276,400
EARLIER FILING DATE: 1999-03-25
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: METHODS OF USE OF A NOVEL LYSYL OXIDASE-RELATED PROTEIN FILE REFERENCE: MNI-073CP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Khodadoust, Mehran et al.
                                 FILE REFERENCE: MNI-073
CURRENT APPLICATION NUMBER: US/09/702,572
CURRENT FILING DATE: 2000-10-31
                                                                                    TITLE OF INVENTION: NOVEL MSP-18 PROTEIN AND NUCLEIC ACID MOLECULES AND TITLE OF INVENTION: USES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.0
PRIOR APPLICATION NUMBER: 09/
PRIOR FILING DATE: 1999-03-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 8.4%;
Local Similarity 68.1%;
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                                                                                                                                                                                                                                                                             2820 TTTTACTATAATATGTTGCTAGGTATAATTTTATTTATATAAAAAAGTGTTTCTGTGATT 2879
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                                                                                                                                                                                                                                                                                                                                                                                           444 TTTTGTAATTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTC 503
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                                                                                                                                                                                                                                                                                                                                                                                                                                    62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (143)..(2401)
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4; Length 2920;
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PRIOR FILING DATE:

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; LOCATION: (143)..(2401)
US-09-702-572-1
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                                                                     Query Match
Best Local :
                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local :
                                                                                                                                                                                            TELEPHONE: (206) 622-490
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 1
                                                                                                                                                                                                                                      CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 2101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Relea
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/O
FILING DATE: 07-0CT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Mitcham, Jennifer L.

TITLE OF INVENTION: COMPOUNDS AND METHODS FOR IMMUNOTHERAPY
TITLE OF INVENTION: AND IMMUNODIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                     Local Similarity
                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0
                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 8.4%;
Local Similarity 68.1%;
les 62; Conservative
441 AAGTTTTGTAATTTTATATTTAGTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTA 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2880 CTTCAGAAAAAAAAAAAAAAAAAAAAAAA 2910
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            444 TTTTGTAATTTTATATTTACTTTTAGTTTGGATACTAAGTATTAAACATATTTCTGTATTC 503
                                                    63; Conservative
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                                                                                                                                                                                  333 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08946026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6300 Columbia Center, 701 Fifth Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reed, Steven G
                                                                                                                                    linear
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                                                                                                                                                single
                                                                 8.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                         Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                           US/08/946,026
                                           Score 44.4; DB 3;
Pred. No. 0.014;
0; Mismatches 31;
                                                                                                                                                                                                                                                                              210121.424C1
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                                                                             Length 333;
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US-08-557-309B-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B
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                                                                                                                                                                                              Patent No. 6054135
GENERAL INFORMATION:
                                                                                                                                                                                                                        Sequence 22, Approximately 100 c. 100
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APPLICANT: Reed, St
APPLICANT: Skeiky,
APPLICANT: Lodes,
APPLICANT: Houghto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22, Application US/08557309B Patent No. 5916572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (206) 682-603
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                 APPLICANT:
                         APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DETECTION AND
NUMBER OF SEQUENCES:
                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION: NAME: Maki, David J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE DETECTION AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 14 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/557,309B
FILING DATE: 14-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 98104-7092
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   544 CAAAAAAAAAAAAAAAAAAAAAAAA 572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 446 TIGIAATITTATATTACTITTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTT 505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        506 CCAAAAAAAAAAAAAAAAAAAAAAAA 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         501 TTCTTCCAAAAAAAAAAAAAAAAAAAAAAAAA 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            293 СТТТТАGААААААААААААААААААААААА 326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       233 AACTTTGGAATATATGTTTGCCAATCTCCTTGTTTTCTAATGAATAAATGTTTTTATATA 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61;
                                                                                                                                                                                                                                                                      Application US/08834306
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                                                                                             Skeiky, Yasir A.W. Lodes, Michael J.
                                                                                                                                                                     Reed, Steven G.
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Lodes, Michael J.
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65
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                            PREVENTION OF
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STATE: washing--
COUNTRY: USA
ZIP: 98104-7092
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/834,306
FILING DATE: 15-APR-1997
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J
REGISTRATION NUMBER: 210121.422C1
REFERENCE/DOCKET NUMBER: 210121.422C1
REFERENCE/DOCKET NUMBER: 210121.422C1
TELEPHONE: (206) 622-4900
TELEPAX: (206) 622-4900
TELEPAX: (206) 622-4900
TELEPAX: (206) 622-6031
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 581 base pairs
TYPE: nucleic acid
STRANDEDNESS: Single
TOPOLOGY: linear
US-08-834-306-22

8.38; Score 44.2; DB 3;
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Search completed: June 25, 2003, 11:55:46 Job time: 68 secs
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ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 6300 Colum
CITY: Seattle
STATE: Washington
                                                                                         446 TIGTAATTITATATTACTTTITAGTTIGATACTAAGTAITAAACATATTITCTGTATTCTT 505
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                        /cgn2_6/piodata/1/pubpna/USO/_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/USO6_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/USO6_PUBCOMB.seq:*
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/cgn2_6/ptodata/1/pubpna/USO8_NEW_FUB.seq:*
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/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
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/cgn2_6/ptcodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptcodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptcodata/1/pubpna/US60_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

19	18	17	16	15	14	13	12	11	10	9	8	7	6	<sub>5</sub>	4	ω	Ν	H	Result No.
335	335	335	335	335	335	335	335	335	335	335	335	404	425	460.2	478.6	486.6	497.2	534	Score
62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	75.7	79.6	86.2	89.6	91.1	93.1	100.0	Query Match
663	663	663	663	663	663	663	663	663	663	663	663	422	439	512	506	504	558	534	Query Match Length
9	9	9	9	9	9	9	9	9	9	9	9	9	9	9	9	9	9	10	DB
US-09-978-824-189	US-10-017-081A-189	US-09-978-585A-189	US-09-978-564A-189	US-09-978-403A-189	US-09-978-191A-189	US-09-978-608A-189	US-09-978-189-189	US-09-999-832A-189	US-09-978-192A-189	US-09-978-697-189	US-09-978-295A-189	US-09-918-995-6534	US-09-822-846-397	US-09-918-995-19489	US-09-809-391-252	US-09-809-391-98	US-10-106-698-1927	us-09-801-115-1	ID
Sequence 189, App	•	Sequence 189, App	•	Sequence 189, App	•	•	Sequence 189, App	Sequence 189, App	•	•	Sequence 189, App	6534,	Sequence 397, App	36	Sequence 252, App	Sequence 98, Appl	Sequence 1927, Ap	Sequence 1, Appli	Description

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JS-09-978-189-191	JS-09-999-832A-191	-09	US-09-978-697-191	-09-	S-09	US-09-954-531-450	-09-	-09-	US-09-801-115-3	-09-	-09	US-10-166-709A-189	-09-	JS-09-978-187B-18	JS-09-978-757A-18	US-09-999-830A-18	JS-10-016-177A-18	JS-10-013-929A	US-09-978-193A	US-10-013-921A-18	-09-	US-09-918-585A-189	-10-167-749-189	:-09-999-833A-189	
-978	-999	-978	-978	978	88-	954	954	954	9-80	.871	822	.166	.978	978	-978	999	016	013	978	013	.978	·918	167	.999	
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## ALIGNMENTS

RESULT 1 US-09-801-115-1

GENERAL INFORMATION:

Ma, D. Han, W.

APPLICANT:

APPLICANT: APPLICANT: Sequence 1, Application US/09801115 Patent No. US20020001828A1

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                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 10776-003-999
CURRENT APPLICATION NUMBER: US/09/801,115
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: PCT/CN00/00026
PRIOR FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 92107284.7
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                              SEQ ID NO 1
LENGTH: 534
                                                                                                                                                                                             Matches 534; Conservative
                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: CHEMOKINE LIKE FACTOR(CKLF) WITH CHEMOTACTIC AND HEMATOPOIET TITLE OF INVENTION: STIMULATING ACTIVITIES FILE REFERENCE: 10776-003-999
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                                                                                                                                                                                           y Match 100.0%; Score 534; DB 10; Local Similarity 100.0%; Pred. No. 1.2e-124; hes 534; Conservative 0; Mismatches 0;
121 GGAAAGTGCTGCTGCGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACA 180
                                                          1 GTTCCCAATCTGAAGTGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCG
                                                                                                                 1 GTTCCCAATCTGAAGTGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCG
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Song, Q.
D1, C.
Huang, J.
Tang, G.
Chen, G.
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RESULT 2
US-10-106-698-1927
; Sequence 1927, Application US/10106698
; Publication No. US20030109690A1
; GENERAL INFORMATION:
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; LOCATION: (514)..(514)
; OTHER INFORMATION: n equals a,t,g,
US-10-106-698-1927
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PRIOR FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 8564
SOFTWARE: Patentin Ver. 3.0
SEQ ID NO 1927
LENGTH: 558
                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                       Query Match
Best Local 9
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CURRENT APPLICATION NUMBER: US/10/106,698
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: PCT/US00/26524
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US 60/157,137
PRIOR FILING DATE: 1999-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ruben et al.
TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypeptide
                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                   131
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                                                                                                          76 TCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCT
                                                                                                                                                       11 TGAAGYSGGAGCTGGCGAGAAGTAGGGGAGGGGGTGCTCCGCCGCGGTGGCGGTTGCTA
                                                                                            71 TCGCTTCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGC
                                                                                                                                                                                         16
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                                                TGGGTCTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTT
                                                                                                                                                                                    TGAAGCCGAGCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTA
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                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                     93.1%;
98.4%;
                                                                                                                                                                                                                   Score 497.2; DB 9;
Pred. No. 2.2e-115;
2; Mismatches 6;
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; ORGANISM: Homo
US-09-809-391-98
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US-09-809-391-98
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                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. SEQ ID NO 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ruben et al.
TITLE OF INVENTION: 186 Human Secreted pro
FILE REFERENCE: PZ002P2
CURRENT APPLICATION NUMBER: US/09/809,391
CURRENT FILING DATE: 2001-03-16
Prior application data removed - consult I
NUMBER OF SEQ ID NOS: 761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 504
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                GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC.
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                                             GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC
                                                                                                                         TCATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTG
                                                                                                                                                                                      TGAAAGGCCACGTGAAGATGCTGCGGGTTGGATATTATCAACTCACTGGTAACAACAGTAT
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Conservative
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99.0%;
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Pred. No. 9.8e-113;
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NUMBER OF SEQ ID NOS: 761
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 252
LENGTH: 506
TYPE: DNA
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GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2
GIDER REFERENCE: PZ002P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 484; Conserv
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CURRENT FILING DATE: 2001-03-16
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RESULT 5 US-09-918-995-19489

RESULT 6 US-09-822-846-397

Sequence 397, Application US/09822846 Publication No. US20030027139A1 GENERAL INFORMATION:

APPLICANT: Jacobs, Kenneth APPLICANT: McCoy, John M.

APPLICANT: APPLICANT: APPLICANT:

LaVallie, Edward R. Collins-Racie, Lisa Evans, Cheryl Merberg, David Treacy, Maurice Agostino, Michael J.

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APPLICANT:

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NAME/KEY: misc_feature; LOCATION: (1)...(512); OTHER INFORMATION: n = US-09-918-995-19489
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LENGTH: 512
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Best Local Similarity
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TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CURRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
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PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
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Sequence 6534, Application US/09918995
Publication No. US20030073623A1
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-756
CUGRENT APPLICATION NUMBER: US/09/918,995
CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING NOVEL SECRETED PROTEINS
FILE REFERENCE: GIN 6400
CURRENT APPLICATION NUMBER: US/09/822,846
CURRENT FILING DATE: 2001-03-29
PRIOR APPLICATION NUMBER: 60/195,605
PRIOR APPLICATION NUMBER: 60/195,605
NUMBER OF SEQ ID NOS: 629
SOFTWARE: Patentin Ver. 2.0
150/THARE: Patentin Ver. 2.0
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ORGANISM: Homo sapiens
-09-822-846-397
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Graham, James R.
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Bowman, Michael R.
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Pred. No. 2.8e-97;
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PRIOR FILING DATE: 1999-01-20
NUMBER OF SEQ ID NOS: 38054
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 6534
LENGTH: 422
TYPE: DNA
ORGANISM: Homo sapiens
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                                                  GACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAG
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US-09-978-295A-189 GENERAL Sequence 189, Application US/09978295A Patent No. US20020156006A1 APPLICANT: APPLICANT: APPLICANT: APPLICANT:
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APPLICANT: APPLICANT: Ashkenazi, Avi APPLICANT: Baker Kevin P. APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: INFORMATION: Fong, Successions Gao, Wei-Qiang Pan, Kuo, Tumas, Daniel Roy, Margaret Ann Shelton, David L. Stewart, Timothy ! Paoni, Kljavin, Ivar J. Gurney, Austin L. Hillan, Kenneth J Godowski, Paul J. Grimaldi, J. Christopher Goddard, Audrey Gerritsen, Mary E. Gerber, Hanspeter Ferrara, Napoleon Filvaroff, Ellen Eaton, Dan Desnoyers, Luc Botstein, David Napier, Mary A. James; Sophia S. Nicholas F.

60/081195

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APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2630P1C11
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FILING DATE: 1998-04-08
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OR FILING DATE: 1998-05-15
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APPLICATION NUMBER: 60/085323
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FILING DATE: 1998-05-07
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PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/066364
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PRIOR APPLICATION NUMBER: 60/077450
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PRIOR APPLICATION NUMBER: 60/064249
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CURRENT FILING DATE: 2001-10-16
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                           APPLICATION NUMBER: 60/078936 FILING DATE: 1998-03-20
                                                           APPLICATION NUMBER: 60/078004
FILING DATE: 1998-03-13
APPLICATION NUMBER: 60/078886
FILING DATE: 1998-03-20
                                                                                                                                   APPLICATION NUMBER: 60/077791
                                                                                                                                                                                                   APPLICATION NUMBER: 60/077641
             APPLICATION NUMBER: 60/078910
                                                                                                                          FILING DATE: 1998-03-1
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Williams, P. Mickey
Wood, William I.
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Stewart, Timothy A.
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Grimaldi, J. Christopher
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Hillan, Kenneth J
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Desnoyers, Luc
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DR FILING DATE: 1998-03-31
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FILING DATE: 1998-04-01
APPLICATION NUMBER: 60/080334
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OR FILING DATE: 1998-5-07
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Query Match Best Local S Matches 504 504; 22 Similarity Conservative 62.78; 76.08; 0; Score 335; DB 9; Pred. No. 1.6e-74; Mismatches Length 663; Indels 159; Gaps

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82 CGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGGGTC 141

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RESULT 10
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; Sequence 189, Application US/09978192A
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APPLICANT: Baker Ke
APPLICANT: Botstein
APPLICANT: Desnoyer
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                                                                                                             Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
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Eaton, Dan
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                                                                   Napier,
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    Margaret Ann
con, David L.
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                                                                                Sophia S.
                                   Nicholas F.
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PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/064249
PRIOR FILING DATE: 1997-11-03
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-3
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PRIOR FILING DATE: 1997-11-21
PRIOR APPLICATION NUMBER: 60/07450
PRIOR APPLICATION NUMBER: 60/077652
PRIOR FILING DATE: 1998-03-10
PRIOR APPLICATION NUMBER: 60/077632
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APPLICANT: Williams, P. Mickey
APPLICANT: William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2630P1C9
CURRENT APPLICATION NUMBER: US/09/978,192A
CURRENT FILING DATE: 2001-10-15
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DR APPLICATION NUMBER: 60/081203
DR FILING DATE: 1998-04-09
DR APPLICATION NUMBER: 60/081229
DR FILING DATE: 1998-04-09
DR APPLICATION NUMBER: 60/081955
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OR APPLICATION NUMBER: 60/084639
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OR APPLICATION NUMBER: 60/084640
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APPLICATION NUMBER: 60/081071
FILING DATE: 1998-04-08
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APPLICATION NUMBER: 60/081049
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FILING DATE: 1998-04-30
APPLICATION NUMBER: 60/084366
FILING DATE: 1998-05-05
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APPLICATION NUMBER: 60/083559
FILING DATE: 1998-04-29
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FILING DATE: 1998-04-29
APPLICATION NUMBER: 60/083554
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APPLICATION NUMBER: 60/083499
FILING DATE: 1998-04-29
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FILING DATE: 1998-04-22
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APPLICATION NUMBER: 60/
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APPLICATION NUMBER: 60/085323
FILING DATE: 1998-05-13
APPLICATION NUMBER: 60/085582
FILING DATE: 1998-05-15
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FILING DATE: 1998-05-07
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APPLICATION NUMBER: 60/085579
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CURRENT FILING DATE: 2001-10-24
PRIOR APPLICATION NUMBER: 09/91858
PRIOR FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: 60/062250
PRIOR TILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/064249
PRIOR APPLICATION NUMBER: 60/065311
PRIOR PILING DATE: 1997-11-03
PRIOR APPLICATION NUMBER: 60/065311
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APPLICATION NUMBER: 60/077632
FILING DATE: 1998-03-11
APPLICATION NUMBER: 60/077641
FILING DATE: 1998-03-11
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FILING DATE: 1997-11-21
APPLICATION NUMBER: 60/077450
FILING DATE: 1998-03-10
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FILING DATE: 1998-03-11
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Shelton, David L.
Stewart, Timothy A.
Tumas, Daniel
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Grimaldi, J. Christopher
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Hillan, Kenneth J
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PRIOR FILING DATE: 1998-03-20
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DR FILING DATE: 1998-04-15
DR APPLICATION NUMBER: 60/081819
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                                                          APPLICATION NUMBER: 6: FILING DATE: 1998-04-
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APPLICATION NUMBER: 60/081955
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FILING DATE: 1998-04-09
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APPLICATION NUMBER: 60/082796
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APPLICATION NUMBER: 60/080165
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Baker Kevin P
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Roy, Mar
Shelton,
                                               Pan,
                                                                                                                Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
                                                                                                                                                                                                                                                                    Filvaroff,
                                                                                                                                                                                                                   Gerber, Hanspeter
                                                                                                                                                                                                                                                                                                                     Desnoyers, Luc
                                                                                                                                                                                                                                                                                                                                         Botstein, David
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o. US20030004102A1
                                 Paoni,
                                                                 Napier, Mary A.
                                                                                  Kuo,
                                                                                                Kljavin,
                                                                                                                                                                                     Goddard, Audrey
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                                                                                                                                                                                                                                                                                       Ferrara,
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                                                                             Sophia S.
                                                   James;
               Margaret
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                                   Nicholas F.
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 David
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CURRENT APPLICATION NUMBER: US/09/978,189
CURRENT FILING DATE: 2001-10-15
PRIOR APPLICATION NUMBER: 09/918585
PRIOR FILLWG DATE: 2001-07-30
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILLWG DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/064249
PRIOR FILLWG DATE: 1997-11-03
PRIOR FILLWG DATE: 1997-11-03
PRIOR APPLICATION NUMBER: 60/065311
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILLING DATE: 1997-11-13
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OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/078936
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OR APPLICATION NUMBER: 60/078910
OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/078939
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OR APPLICATION NUMBER: 60/079294
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R APPLICATION NUMBER: 60/080105

DR FILING DATE: 1998-03-31

DR APPLICATION NUMBER: 60/080107

DR FILING DATE: 1998-03-31

DR APPLICATION NUMBER: 60/080165

DR FILING DATE: 1998-03-31
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OR FILING DATE: 1998-03-27
OR APPLICATION NUMBER: 60/079689
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CGGTCCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAAGAAGTTTTGTAATTTTATATTAC 462
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CURRENT FILING DATE: 2001-10-16
NUMBER OF SEQ ID NOS: 624
Prior Application removed - See File Wrapper or Palm
SEQ ID NO 189
LENGTH: 663
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Williams, P. Mickey
Wood, William I.
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Shelton, David L.
Stewart, Timothy A.
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Hillan, Kenneth J
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Grimaldi, J. Christopher
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Gao, Wei-Qiang
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Pred. No. 1.6e-74;
0; Mismatches 0;
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; APPLICANT: WOOD, WILLIAM I. ; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic; TITLE OF INVENTION: Acids Encoding the Same ; FILE REFERENCE: P2630P1C4 ; CURRENT APPLICATION NUMBER: US/09/978,191A ; CURRENT FILING DATE: 2001-10-15 ; PRIOR APPLICATION NUMBER: 09/918585 ; PRIOR FILING DATE: 2001-07-30	Shelton, David Stewart, Timor Tumas, Daniel Williams, P. 1		; APPLICANT: Goddard, Audrey ; APPLICANT: Goddwski, Paul J. ; APPLICANT: Grimaldi, J. Christopher ; APPLICANT: Gurney, Austin L. ; APPLICANT: Hillan, Kenneth J ; APPLICANT: Kliavin, Ivar J.	: Filvarof : Fong, Sh : Gao, Wei : Gerber, 1 : Gerritse	<pre>: Botstein, D : Desnoyers, : Eaton, Dan : Ferrara, Na</pre>	plicati US2003( ION: enazi, er Kevi	DD 001 AAA 003 RESULT 14 US-00-978-191A-189	523 AAA 	Qy 463 TTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTTCCAAAAAAAA	Qy 403 CGGTCCTTACCAGAAAAAAGCCTGTGCATGAAAAAAAAAA	Qy 343 AGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTTCTGTTCAATCCCAG 402	Qy 283 GGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGAC 342	Qy 230GATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTGTT 282	301 ATTTTTCATACTTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCC	Db 241 CATCGCACAAGCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTT 300 Ov 230	ОУ 230 229
PRIOR FILING DATE: 1998-04-08  PRIOR APPLICATION NUMBER: 60/081071  PRIOR APPLICATION NUMBER: 60/081195  PRIOR FILING DATE: 1998-04-08  PRIOR FILING DATE: 1998-04-08  PRIOR APPLICATION NUMBER: 60/081203  PRIOR APPLICATION NUMBER: 60/081229  PRIOR APPLICATION NUMBER: 60/081229	CATION PERSONS AND ADDRESS OF DATE:	PRIOR FILING DATE: 1998-04-01 ; PRIOR FILING DATE: 1998-04-01 ; PRIOR APPLICATION NUMBER: 60/080333 ; PRIOR FILING DATE: 1998-04-01 ; PRIOR FILING DATE: 1998-04-01	APPLICATION   FILING DATE: APPLICATION   FILING DATE: APPLICATION   APPLICATION	FILING DATE: 1998-0: APPLICATION NUMBER: FILING DATE: 1998-0: APPLICATION NUMBER: FILING DATE: 1998-0:	FILING DATE: 1998-0: APPLICATION NUMBER: FILING DATE: 1998-0: APPLICATION NUMBER:	PRIOR FILLING DATE: 1996-03-27  PRIOR APPLICATION NUMBER: 60/07963  PRIOR FILING DATE: 1998-03-27  PRIOR FILING DATE: 1998-03-27  PRIOR FILING DATE: 1998-03-27  PRIOR APPLICATION NUMBER: 60/079786	FILING DATE APPLICATION FILING DATE APPLICATION APPLICATION		; PRIOR APPLICATION NUMBER: 60/0/8910 ; PRIOR FILING DATE: 1998-03-20 ; PRIOR FILING DATE: 1998-03-20 ; PRIOR FILING DATE: 1998-03-20			FILING DATE: 1998-03-11 APPLICATION NUMBER: 60 FILING DATE: 1998-03-11	FILING DATE: 1998-0 APPLICATION NUMBER: FILING DATE: 1998-0 ADDITCATION NUMBER	PRIOR FILING DATE: 1997-11-21	FILING DATE: 1997: 11-03 APPLICATION NUMBER: 60, ETITING DATE: 1997-11-13	; PRIOR APPLICATION NUMBER: 60/062250 ; PRIOR FILING DATE: 1997-10-17

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PRIOR FILLING DATE: 1998-04-09
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RESULT 15 US-09-978-403A-189

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Sequence 189, Application US/09978403A Publication No. US20030050240A1 GENERAL INFORMATION:
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APPLICANT: Baker Kevin P
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APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
TILE REFERENCE: P2630P1C17
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RIOR FILING DATE: 1997-10-17
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R APPLICATION NUMBER: 60/07
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Botstein, David
Desnoyers, Luc
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Filvaroff,
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Shelton, David L.
Stewart, Timothy A.
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Grimaldi, J. Christopher
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Hillan, Kenneth J
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Gerritsen, Mary E
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Ef, Ellen
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APPLICATION NUMBER: 60/085338
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NA 52:	TTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTTCCAAAAAAAA	463	Qy
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C 342	GGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGAGGGGTGTTTGCACTTGTGAC	283	Qy
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C 360	ATTTTTCATACTTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCC	301	Db
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T 300	CATCGCACAAGCCCCTGAACCATATATTGTTATCACTGGATTTGAAGTCACCGTTATCTT	241	рь
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## SUMMARIES

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482.4	485.4	487	491.8	492.4	505.8	Score
90.3	90.9	91.2	92.1	92.2	94.7	
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AI349474	BI561562	AI371387	AI263261	BF388158	AA644621	BE737159	BQ013860	AA987264	AA477156	N89912	AF151058	AI265924	BE875161	AA305052	W38899	W52820	AII41284	AA989129	AA911088	AI989739	AW183193	AA516431	AI989747	BM553628	AI826623	BF215121	BG532587	AV759888	BG613984	AI128804	AA455042	AA429945	AV754613	AI743235	AI078580	BF203359	BF109912	BF691818
AI3494/4 qp/2IU/.X	BI561562 603256264	AI371387 go84b02.x	AI263261 qp62b12.x	BF388158 UI-R-CA1-	AA644621 af74c11.r	BE737159 601305091	BQ013860 UI-1-BClp	AA987264 oslla06.s	AA477156 zu41b02.r	N89912 zb22g09.sl	AF151058 Homo sapi	AI265924 qx68d08.x	BE875161 601488641	AA305052 EST176055	W38899 zb32d04.rl	W52820 zc55c06.rl	AI141284 qa52h10.s	AA989129 or75h10.s	AA911088 ok67e01.s	AI989739 wu2lel0.x	AW183193 xj67f11.x	AA516431 ne58a03.s	AI989747 wu21f09.x	BM553628 AGENCOURT	AI826623 wk35d10.x	BF215121 601846829	BG532587 602562152	AV759888 AV759888	BG613984 602641513	AI128804 qa94b08.s	AA455042 aa04a07.s	AA429945 zw67f10.s	AV754613 AV754613	wg9(	AI078580 oz37h05.x	BF203359 601865968	BF109912 7171f03.x	60224778

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## ALIGNMENTS

FEATURES	COMMENT	AUTHORS	REFERENCE	Circuit	SOURCE	VERSION KEYWORDS	ACCESSION	RESULT 1 BG705303 LOCUS DEFINITION
Email: cgapbs remail.nin.gov Tissue Procurement: Miklos Palkovits, M.D., Ph.D. CDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki Toshiyuki and Piero Carninci (RIKEN) CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLAM10725 row: o column: 01 High quality sequence stop: 521. Location/Qualifiers	Contact: Robert Strausberg, Ph.D.	NATH-MGC http://mgc.nci.nin.gov/. National Institutes of Health, Mammalian Gene Collection (MGC)		Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.	human. Homo sapiens	EST.	mRNA sequence. BG705303	BG705303 524 bp mRNA linear EST 07-MAY-2001 602687808F1 NIH_MGC_95 Homo sapiens cDNA clone IMAGE:4820568 5',

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              Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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     (bases 1 to 962)
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/db_xref="taxon:9606"
/clone="IMAGE:4820568"
/clone_lib="NIH_MGC_95"
/tissue_type="hippocampus"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhOI (gtcga); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.5 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library. "a 113 c 138 g 131 t
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Pred. No. 8.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               89 CCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGCGGTCTGCAGAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     http://image.llnl.gov
Plate: LLAM10288 row: k column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                      AAAAAAAAAAAAAAAAAAAAAAAA 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                           TAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATFCTTCCA
                                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                       TAATTTTATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTTCCA
                                                                                                                                                                                                           TTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGGAAGCTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="adenocarcinoma, cell line"
/lab_host="DHIOB (phage resistant)"
/note="Organ: liver; Vector: pcMv-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally; oligo-dT primed.
Average insert size 1.7 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
a 199 c 288 g 141 t
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/clone_lib="NIH_MGC_90"
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Pred. No. 5e
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                                                                                                                                                                                                                                                                              435
                                                                                                                                                                                                                148 CGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGG
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UI-R-CA1-bjb-b-12-0-UI.sl UI-R-CA1 Rattus norvegicus cDNA clone
UI-R-CA1-bjb-b-12-0-UI 3', mRNA sequence.
BF399486
                                                                                                                                                                                                                                                                                                                                            88 ACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGCTGGGTCTGCAGA 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          strand cDNA and therefore this may represent a bonafide poly A tail. cDNA Library Preparation: M.B. Soares Lab Clone distribution: clones will be available through Research Genetics (www.resgen.com)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: msoares@blue.weeg.uiowa.edu
The sequence contained an oligo-dT track that was present in oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide that was used to prime the synthesis of first oligonucleotide the synthesis of the synthesis oligonucleotide that was used to prime the synthesis of the synthesis oligonucleotide the synthesis of the synthesis oligonucleotide that was used to be something the synthesis of the synthesis oligonucleotide th
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Program for Rat Gene Discovery and Mapping
University of Iowa
451 Eckstein Medical Research Building Iowa
Tel: 319 335 8250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Rattus norvegicus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                           CCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCT
                                                                                                                                                                                                                                                                                 ACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGAAAGTGCTGCTGGTGGGTCTGCAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                    GGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCTTCGCAGA
                                CCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCT
                                                                                                                                                         CGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: pT/T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; The UI-R-CAl library is a subtracted library derived from the following tissues: thalamus, cerebellum, hypothalamus, medulla, pons, midbrain, cerebral cortex, corpus striatum, testis, and hippocampus. For a detailed description of the library from which this clone was derived, please visit our web site at ratest eng.ulowa.edu. The subtraction has been previously described in (Bonaldo, Lennon and Soares, denome Research 6:791-806, 1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TAG_SEQ=None
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /strain="Sprague-Dawley"
/db_xref="taxon:10116"
/clone="UI-R-CA1-bjb-b-12-0-UI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="UI-R-CA1"
/lab_host="DH10B (Life Technologies)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Rattus norvegicus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 491.8; DB 1
Pred. No. 9.9e-64;
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCCTCC
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                                                                                                                           a
                                                                                                                               Library.
                                                                                                                                                                                                                /note="organ: prostate; vector: pDNR-LIB (Clontech); Site_1: Sfil (ggccgctcggcc); Site_2: Sfil (ggccgtctggcc); Site_2: Sfil (ggccgttattggcc); Double-stranded cDNA was prepared from cell line RNA.
5' and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCGATTATGGCC-3' and 3' adaptor sequence: 5'-CACGGCCGATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGAGGCGGCGCATGTG-dT[30]BN-3'
(where B = A, C, or G and N = A, C, G, or T). Average insert size 1.5 kb (range 0.9-4.0 kb). 14/15 colonies contained inserts by pCR. This library was enriched for full-length clones and was constructed by Clontech
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="NIH_MGC_60"
/tissue_type="adenocarcinoma"
/lab_host="DH10B (T1 phage-resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4713305"
                                                                                                                                                                                         Laboratories (Palo Alto, CA). Note: this is a NIH_MGC
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                                                                                                                              found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
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                                                                                                    quality sequence stop: 486.
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/organism-"Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5576339"
/clone_lib-"NIH_MGC_92"
/tissue_type="embryonal carcinoma,
                                                                                      ocation/Qualifiers
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Best Local S
Matches 486
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                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                        NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                    mRNA sequence.
BG031757
BG031757.1 GI:12422364
                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 534)
                                                                                                                                                                                                                                                                          BG031757 534 bp m
602299819F1 NIH_MGC_87 Homo sapiens
                                                                                                                                                                                            Homo sapiens
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CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           ATATTACTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTATTCTTCCAAAAAAA 515
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/note="Organ: testis; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: Sali; Cloned unidirectionally; oliqo-dT primed.
Average insert size 2.5 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
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Pred. No. 9e
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RESULT 7
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BF691818.1 GI:11977226
                                                                                                                                                602247787F1 NIH_MGC_62 Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                     Homo sapiens
                                                        human.
                                                                                                                              mRNA sequence.
                                                                                                                                                                  BF691818
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                               CCCACAAAAAAAAAAAAAAAAAAAAA 508
                                                                                                                                                                                                                                                                                                  CAAAAAAAAAAAAAAAAAAAAAAAA 534
                                                                                                                                                                                                                                                                                                                                      AATTTTATATTACTTTTAGTTTGATACTAAGTTATTTAAAACATATTTCTGTATTCTTT
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/clone_lib="NIH_MGC_893"
/clone_lib="NIH_MGC_893"
/clone_lib="NIH_MGC_893"
/tissue_type="mammary adenocarcinoma, cell line"
/lab_host="DH10B (phage-resistant)", Sporm5; Site_1: NotI, note="Organ: breast; Vector: pcWV-Sporm5; Site_1: NotI; Site_2: Sall; cloned unidirectionally; oligo-dT primed. Average insert size 1.383 kb. Library enriched for full-length clones and constructed by Life Technologies. Note: this is a NHL_MGC_Library."

108 c 132 g 131 t
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/db_xref="taxon:9606"
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98.2%;
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Pred. No. 2.3e-62;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLCM1200 row: m column: 21
High quality sequence stop: 483.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           National Institutes of Health, Mammalian Gene
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NIH-MGC http://mgc.nci.nih.gov/.
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                                                                                                           ATGCTCATCGTATCTGTGTTGGCACTGATACCAGAAACCACAACATTGACAGTTGGTGGA
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                                                                                                                                                                GGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACCGG
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                                                    AGTTTTGTAATTTTATATTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTAT 501
                                                                                       AAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCTGTGCATGAAACAAAACAG
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/lab_host="DH10B (TI phage resistant)"
/note="Organ: skin; Vector: pDNR-LIB (Clontech); Site_1:
/note="Organ: skin; Vector: SFII (ggccattatggcc);
SfII (ggccgctcggcc); Site_2: SFII (ggccattatggcc);
Double-stranded cDNA was prepared from cell line RNA. 5'
and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
sequence: 5'-ATTCTAAAGGCCGACATCATG-dT(30)BN-3'
(where B = A, C, Or G and N = A, C, G, Or T). Average
insert size 1.75 kb (range 0.9-4.0 kb). 15/15 colonies
contained inserts by PCR. This library was enriched for
full-length clones and was constructed by Clontech
Laboratories (Palo Alto, CA)."
39 a 119 c 138 g 133 t
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/clone_lib="NIH_MGC_62"
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RESULT 8
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468 bp mRNA linear EST 20-OCT-200: 7171f03.xl Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone limage:3526805 3' similar to TR:Q9UI41 Q9UI41 CHEMOKINE-LIKE FACTOR 1.[1]; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Seq primer: -400P from Gibco.
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BF109912.1 GI:10939602
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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National Cancer Institute, Cancer Genome Anat
   GCTGGATATTATCAACTCACTGGTAACAACAGTATTCATGCTCATCGTATCTGTTTGGC
                            GCTGGATATTATCAACTCACTGGTAACAACACATATTCATGCTCATCGTATCTGTGTTGGC 285
                                                                                                                                                  GCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCCACGTGAAGATGCTGCG
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116 c 102 g 123 t
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
/lab_host="DH10B"
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100.0%;
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Pred. No. 3.4e-60;
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                                                                                                                                                  25 GCTGGGCGAGAAGTAGGGGAGGGCGGTGCTCCGCCGCGGTGGCGGTTGCTATCGCTTCGC
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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601865968F1 NIH_MGC_17 Homo
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Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
Plate: LLCM965 row: h column: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1999)
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318
                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="rhabdomyosarcoma"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: muscle; Vector: pOTB7; Site_1: EcoRI;
Site_2: Xho1; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCAGGAG(G). Size-selected >500bp
                                                                                                                                                                                                                                                                                                                                                          for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."

203 c 257 g 152 t
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/clone_lib="NIH_MGC_17"
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                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL ;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anai
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1 (bases 1 to 452)
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Seq primer: -40m13 fwd. ET from Amersham
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                                                                                   /note-"Organ: mixed (see below); Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and so circles
                  were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools
                                                                                                                                                                                                /clone_lib="Soares_NhHMPu_S1"
/tissue_type="Pooled human me
pregnant uterus"
     consisted of I.M.A.G.E. clones 260232-265223
                                                                                                                                                                                                                                                      /clone="IMAGE:1677561"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov This clone is available royalty-free through LLNL; contact the
                                                                                                                                                                                                                                                    Unpublished (1997)
Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 453)
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AI743235.1 GI:5111523
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453 bp mRNA linear EST 20-DEC wg90a02.xl Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone mage:2372330 3', mRNA sequence.
                                                                                                                                                           Seq primer: -40UP from Gibco.
                                                                                                                                                                            IMAGE Consortium (info@image.llnl.gov) for Insert Length: 378 Std Error: 0.00
                                                                                                                                                                                                                                                                                                          National Cancer Institute, Cancer
                                                                                                                                                                                                                                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                         Gene Index
                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2372330"
/clone_lib="Soars_NSF_F8_9W_OT_PA_P_S1"
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110 c 97 g 120 t
/note="Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with
a modified polylinker; Site_1: Not I; Site_2: Eco RI;
                                     /lab_host="DH10B"
                                                                                                                                          location/Qualifiers
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Pred. No. 8.1e-58;
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              Song,H., Peng,Y., Gu,Y., Yang,Y., Gao,G., Xiao,H., Xu,X., Li,N., Qlan,B., Liu,F., Qu,J., Gao,X., Cheng,Z., Xu,Z., Zeng,L., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhong,M., Lu,G., Hu,R., Ye,M., Zhang,Q., Han,Z., Chen,Z. and Chen,J. Homo sapiens TP library cDNA clones Unpublished (2000)
                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 533)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the same 5 libraries. The pools consisted of the following libraries and cloneIDs: Soares NbHSF pool 1: 309384-310919, 333208-325895 Soares Nb2HP pool 1: 145032-147335, 147720-148103, 148872-149255, 15002-150407, 151176-152327 Soares Nb2HF8-9W pool 1: 758280-760583, 772104-774407 Soares NbHRA pool 1: 304776-306311, 320136-322823, 326280-32663 Soares NbHOT pool 1: 733720-726407, 739080-740999 Subtraction by Bento Soares and M. Fatima Bonaldo."
Qinghua Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      libraries were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Equal amounts of plasmid DNA from five normalized
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Pred. No. 8.7e-57;
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e TPAAOAll 5', mRNA sequence.
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                                                     3', mRNA sequence
AA429945
                                                                                     zw67f10.sl Soares_testis_NHT Homo sapiens cDNA clone IMAGE:781291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shanghai Institute of Endocrinology, Rui-Jin Hospital
197 Rui-Jin II Road, Shanghai 200025, P. R. China
Tel: 86-21-64370045(ex.663332)
Fax: 86-21-64743206
Email: mbshl@ms.stn.sh.cn
                                   AA429945.1 GI:2113244
                                                                                                       AA429945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This clone is available at Shanghai Hematology Institute
Shanghai.
Chinese National Human Genome Center at Shanghai
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                             TATTTACCGGAAGCTTCTGTTCAATCCCANCGGTCCTTACCAGAAAAAGCCTGNGCATGA
                                                                                                                                                                                                                                                                                                                                                TATTTACCGGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAAGCCTGTGCATGA 432
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/dev_stage="Adult"
/lab_host="BM25.8"
/note="Vector: pTriplEx2; Site,
112 c 145 g 120 t
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/db_xref="taxon:9606"
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Pred. No. 5.6e-55;
D; Mismatches 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R.
WashU-Merck EST Project 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Wilson RK
Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: est@watson.wustl.edu
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                                                                 AAGTTTTGTAATTTTATATTTACTTTTTAGTTTGATACTAAGTATTTAAACATATTTCTGTA 500
                                                                                                                               GGAAGCTTCTGTTCAATCCCAGCGGTCCTTACCAGAAAAAGCCCTGTGCATGAAAAAAAG
                                                                                                                                                                                                                                          GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC
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                                               AAGTTTTGTAATTTATTATTACTTTTTAGTTTGATACTAAGTATTAAACATATTTCTGTA 13
                                                                                                                                                                                                                     GAGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGACGGGGCCCTTATTTACC
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/db_xref="taxon:9606"
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Pred. No. 1.3e-54;
0; Mismatches 1;
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nes 427; Conserv
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WashU_Merck_EST_Project 1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This clone is available royalty-free through LIML; contact the IMAGE Consortium (infe@image.llnl.gov) for further information. Seq primer: -41ml3 fwd. ET from Amersham High quality sequence stop: 395.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4444 Forest Park Parkway,
                                             TGAAAGGCCACGTGAAGATGCTGCGGCTGGATATTATCAACTCACTGGTAACAACAGTAT 260
                                                                                                                                                        CTGCAGACGCGATAGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTG 200
                                                                                                                                                                                                 TCGCAGAACCTACTCAGGCAGCCAGCTGAGAAGAGTTGAGGGGAAAGTGCTGCTGCTGGGT 368
                                                                                                                                                                                                                          TCGCAGAAACCTACTCAGGCAGCCAGCTGAGAAAGAGTTGAGGGAAAAGTGCTGCTGCTGGGT 140
                         TGAAAGGCCACGTGAAGATGCTGCGCTGGATATTATCAACTCACTGGTAACAACAGTAT 248
                                                                                                               CTGCAGACGCGATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTG
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                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                         reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note="Organ: mixed (see below); Vector: pT/T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NDHM, pregnant uterus NDHPU, and fetal heart NDHH19W) were mixed, and so circles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    were made in vitro. Following HAP purification, the was used as tracer in a subtractive hybridization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="GDB:6043155"
/db_xref="taxon:9606"
/clone="IMAGE:812244"
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IMAGE:812244
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                                                                                                                                                                                                                                                                                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 423)
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                                                            116 a
79.0%; Score 422; DB 9; ilarity 100.0%; Pred. No. 2.3e-53; Conservative 0; Mismatches 0;
                                                                         NbHL19W."
                                                                                                                                                                                                                                              /clone_lib="Soares_fetal_heart_NbHH19W"
                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
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			-	501		141	124	81	184	321	244	261	304	201	364	141

Search completed: June 25, 2003, 11:54:33 Job time: 1427 secs

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Command line parameters:

MODEL-frame+_p2n.model -DEV=xlp

-MODEL-frame+_p2n.model -DEV=xlp

-Q-/Ggn2_1/USPTO_Spool_VS0801115/runat_20062003_141102_13583/app_query.fasta_1.263

-DB-GenEmb1 -QFMT-fastap -SUFFIX-rge -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0

-UNITS-blits -STARF=1 -END=-1 -MATRIX-blosum62 -TRANS-buman40.cdi -LIST=45

-DCCALIGN=200 -THR_SCORE-pot -THR_MAX=100 -THR_INIT=0 -ALIGN=15 -MODE-LOCAL

-OUTFMT-pto -NORM-ext -HEAPSIZE=500 -MINLEN=0 -ALIGN=15 -MODE-LOCAL

-USER-US09801115_0CGN 1_1_3745_0runat_20062003_141102_13583 -NCU=6 -ICPU=3

-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG

-DEV_TIMEDUT-120 -WARN_TIMEDUT-30 -THERADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6

-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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Listing first 45 summaries
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37: em_htg_vrt:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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1 (bases 1 to 515)

dumas milne Edwards, J.B., Bougueleret, L. and Jobert, S.

Complementary dna's encoding proteins with signal peptides
Patent: WO 0100806-A 29 04-JAN-2001;
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/db_xref="GI:12406710"
/translation="MDNVQPKIKHRPFCFSVKGHVKMLRLDIINSLVTTVFMLIVSVL
/translation="MDNVQPKIKHRPFCFSVKGHVKMLRLDIINSLVTTVFMLIVSVL
ALIPETTILTVGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
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US-09-801-115B-2 (1-99) x AF096895 (1-530)
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Query Match:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      148 ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAGGCCAC
                                                                                                                                                        61 AlaLeuValThrAlaValCysCysLeuAlaAspGlyAlaLeuIleTyrArgLysLeuLeu
                                                                                                                                                                                                                                                                                                                              41 ValSerValLeuAlaLeuIleProGluThrThrThrThrLeuThrValGlyGlyGlyValPhe
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2 (bases 1 to 530)
Ban,W.L., Li,Y., Zhang,Y.M., Di,C.H.
Direct Submission
Submitted (03-0CT-1998) Immunology,
Submitted (03-0CT-1998), China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chemotactic activity
Biochem. J. 357 (Pt 1), 127-135 (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Science Center, Xueyuan Road 38, Beijing 100083, China Sequence update by submitter On Sep 7, 2000 this sequence version replaced gi:6288733 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (07-SEP-2000) Immunology, Peking University Health Science Center, Xueyuan Road 38, Beijing 100083, China
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/codon_start=1
/codon_start=1
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/protein_id="AAF06722.1"
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/db_xref="G1:6288734"
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ALIPETTTLTVGGGVFALVTAVCCLANGALIXRKLLFNPSGPYQKKPVHEKKEVL"
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/note="ev-
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                                              IleIleAsnSerLeuValThrThrValPheMetLeuIleValSerValLeuAlaLeuIle 47
                                                                                                         CTTTTATATGTACTCAGACTTGATCGATTAATGAAGTGGTTATTTTGGCCTTTGCTTGAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /product-"transmembrane proteolipid"
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XIVITGFEVTVILFFIILYVLRLDRLMKWLFWPLLDIINSLYTTVFWLIVSVLALIPE
TTTLTVGGGVFFALVTAVCCLADGALIYKKLLFNPSGPYQKKPVHEKKEVL"
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/tissue_type="pancreas"
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/db_xref="taxon:9606"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="GI:12406790"
/db_xref="GI:12406790"
/translation="mdnvQpkikhrpfcFsvkghvkmLrlalfvTsmTffiiAQAPEP
/translation="mdnvQpkikhrpfcFsvkghvkmLrlalfvTsmTtpfillvSLvTtvEnLalfpe
YIVITGFETTLTVLRLDRLMKWLFWPSLDIINSLVTTVFMLLVSVLALIPE
TTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGFYQKKPVHEKKEVL"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Steven Jones, Jennifer Asano, Ian Bosdet, Yaron Butterfield, Susanna Chan, Readman Chiu, Chris Fjell, Erin Garland, Ran Guin, Letticia Hsiao, Martin Krzywinski, Reta Kutsche, Oliver Lee, Soo Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven Ness, Pawan Pandoh, Anna-Liisa Prabhu, Parvaneh Saeedi, Jacqueline Schein, Duane Smailus, Michael Smith, Lorraine Spence, Jeff Stott, Michael Thorne, Miranada Tsai, Natasja van den Bosch, Jill Vardy, George Yang, Scott Zuyderduyn, Marco Marra.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRAL Plate: 13 Row: f Column: 14
This clone was selected for full length sequencing because it in the was selected for full length sequencing because it is the was selected for full length sequencing because it is the was selected for full length sequencing because it is the was selected for full length sequencing because it is the was selected for full length sequencing because it is the way and the way are the way and the way are the w
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Genome Sequence Centre,
BC Cancer Agency, Vancouver, BC, Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-remail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (12-MAR-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 688)
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                                                  /product="Unknown (protein for MGC:10658)"
/protein_id="AAH04380.1"
/db_xref="G1:13325134"
/translation="MDNVQPKIKHRPECESVKGHVKMLRLALTVTSMTFFIIAQAPEP
YIVIIGFEVTYVILEFILLYVLRLDRLMKWLFWPLLDIINSLYTTVFMLIVSVLALIPE
                                  TTTLTVGGGVFALVTAVCCLADGALIYRKLLFNPSGPYQKKPVHEKKEVL"
                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="mGC:10658 IMAGE:3639550"
/tissue_type="Pancreas, adenocarcinoma"
/clone_lib="NIH_MGC_39"
/lab_host="DH10B-R"
                                                                                                                                                                                                                     /codon_start=1
                                                                                                                                                                                                                                                                           note="Vector: pOTB7"
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                                                                                                    submitted (07-SEP-2000) Immunology, Peking University Health Science Center, Xueyuan Road 38, Beijing 100083, China Sequence update by submitter on Sep 7, 2000 this sequence version replaced gi:6630853.
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                                                                                                                                                                                                                                                                                                                                                                                   Submitted (17-MAR-1999) Immunology, Beijing Medical University,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MetAspAsnValGlnProLysIleLysHisArgProPheCysPheSerValLysGlyHis
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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(bases 1 to 689)
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                      /organism="Homo sapiens"
/db_xref="taxon:9606"
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YIVITGEFEVTVILFETILTYVLRLDRLMKMLFWFLLDIINSLVTTVFMLIVSVIALIPE
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/protein_id="AAF19599.1"
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Stanford University School of Medicine
Department of Genetics, M-344, Stanford,
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                       CCTTACCAGAAAAAGCCTGTGCATGAAAAAAAAAAGAAGTTTTG
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Sequence update by submitter
On Sep 7, 2000 this sequence version
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 593)
Han, W.L., Gu, W.F., Li,Y., Zhang,Y.M., Di,C.H., Song,Q.S. and
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US-09-801-115B-2 (1-99) x AF135381
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Submitted (07-SEP-2000) Immunology, Peking University Her
Science Center, Xueyuan Road 38, Beijing 100083, China
Sequence update by submitter
On Sep 7, 2000 this sequence version replaced g1:6630855
Location/Qualifiers
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Han,W.L., Gu,W., Li,Y., Zhang,Y., Song,Q., Di,C. and Ma,D.
Direct Submission
Submitted (17-MAR-1999) Immunology, Beljing Medical University, Xue
Yuan Road, Beljing 100083, China
2 (bases 1 to 434)
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Mammalia; Eutheria;
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                                                     ATGGATAACGTGCAGCCGAAAATAAAACATCGCCCCTTCTGCTTCAGTGTGAAAAGGCCAC
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RKLLFNPSGPYQKKPVHEKKEVL"
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US-09-801-115B-2 (1-99) x AF253064 (1-523)  QY    MetaspasnvalGinProLysIieLysHisArgProPheCysPheSerValLysGlyHis 20	Alignment Scores:  Pred. No.:  9. 14e-26  Alterda No.:  10. 14e-26  10. 15e-26  10. 15e-26	RESULT 11 AP233064 LOCUS AP233064 LOCUS DEFINITION RATUS norvegicus chemokine-like factor 1 (CKLF1) mRNA, complete cds. ACCESSION VERSION RETWORMS SOURCE ORGANISM RATUS norvegicus REFERENCE LOUY, Li, Y., Han, W., Song, Q., Zhang, Y., Di, C. and Ma, D. TITLE JOURNAL Xueyuan Road, Beijing 100083, China REMARK RAMARK

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ACCESSION
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                                                                                                             Young, P.E., Augustus, M., Carter, K.C., Ebner, R., Endress, G., Horrigan, S., Soppet, D.R. and Weaver, Z. Cancer gene determination and therapeutic screening using signature
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Sequence 1296 from Patent WO0194629.
AX330787
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Location/Qualifiers
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Cancer gene determination and therapeutic screening using signature
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.
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Cancer gene determination and therapeutic
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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screening

using signature

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PAT 09-JAN-2002

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1 others

Search completed: June 25, 2003, 13:13:15 Job time: 1316 secs	92 ValHisGiuLysLysGluValLeu	145	 		Db 205 ACATTGACAGTTGGAGGGGGTGTTTGCACTTGTGACAGCAGTATGCTGTCTTGCCGAC 146	Qy 52 ThrLeuThrValGlyGlyGlyValPheAlaLeuValThrAlaValCysCysLeuAlaAsp 71		US-09-801-115B-2 (1-99) x AX408097 (1-207)	Ouery Match: 48.23% Indels: 0 DB: 6 Gaps: 0	imilarity: 97.928	97.928 Matches:	ent Scores: 7.42e-23	CHIGIN	/organism="Homo sapiens" /db_xref="taxon:9606" /note="EMBL/GenBank Accession No. AA338760" BASE COUNT 70 a 43 c 39 g 54 t 1 others	source 1. 207	FEATURES Location/Qualifiers	JOURNAL Patent: WO 0229103-A 744 11-APR-2002;	Gene expression profiles in liver cancer	AUTHORS Alvares, C., Horne, D., Peres-da-Silva, S. and Vocklev, J.G.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. REFERENCE 1	ORGANISM Homo sapiens	SOURCE human.	VERSION AX408097.1 GI:21440802	ON AX408097	N Sequence 744 from Patent WO0229103.	2002 Entert 14-JUN-2002
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